Table 6

ACC. NO.	GI NUMBER
W84560	1395672
W84627	
W84716	1395738 1395825
W84751	1395870
W84774	1395894
W84778	1395898
W84315	1395898 1396014
W85697	1398344
W857C9	1398150
W85764	1398425
W85832	1398262
W85878	1398307
W85881	139831C
W85931	1398520
W86075	1398524
W86124	1398554
W86160	1398639
W86183	1398623
W86196	1398746
W86202	1398642
W86214	1398827
W86282	1398720
W86378	1398196
W86466	1400213
W86653	1400529
W86779	1400507
W86823	1400622
W86832	1400561
W86860	1400589
W86987	1400744
W87281	1401476
W87312	1401377
W87509	1401656
W87522	1401728
W87710	1401764
W87801	1401886
W87823	1401898
W87826	1401901
W87865	1401930
W87891	1401976
W88623	1404173
W88720	1404202
W98737	1404219
W38786	1404477
W88792	1404264
W38823	1404295
W38843	1404340 1404408
W98856	
W88942 W88999	1403848 1403885
W89057	1404082
W89074	1404082
W90036	1404167
W90056	1406077
W9C128	1406118
W90164	1406303
W90224	1406303
1170223	1.002_4

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Table 6

ACC. NO.	GI NUMBER
W90323	1406703
W90364	1406602
W90381	1406138
W90519	1406324
W9C520	1406325
W90529	1406395
W90705	1406651
W90732	1406678
W91890	1424282
W92011	1424374
W92263	1424628
W92278	1424663
W92431	1424815
W92798	1421951
W92964	1422116
W92970	1422141
W93147	1422516
W93315	1422457
W93379	1422501
W93483	1422752
W93500	1422837
W93523	1422644
W93585	1422706
W93638	1422779
W93655	1422845 1422804
W93682	1422851
W93709 W93717	1422859
W93718	1423061
W94009	1423130
W94121	1423244
W94289	1423410
W94316	1423437
W94398	1423529
W94486	1423765
W94591	1423713
W94647	1423829
W95001	1424170
W95050	1424398
W95082	1424200
W95118	1424236
W95279	1425205
W95409	1425324
W95414	1425329
W95757	1425712
W96107	1426014
W96114	1426021
W96134	1426060
W96155	1426061
W96187	1426093
W96189	1426095
W96197	1426103
W96214	1426140
W96216	1426142
W96273	1426180
W96280	1426187
W99251	1432178

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Table 6

ACC. NO.	GI NUMBER
XC0351	28251
XC0497	32130
X00630	11585
MC0632	10627
MC1037	36086
XC163C	28871
X01742 X02422	35324 33344
X02422 X02457	32153
X02530	33917
X02544	24444
X02596	29420
X02761	31396
X02875	35124
X02974	10873
X03559	28939
X04098	28338
K04106	35327
X04125	58114
X04297	28926
X04318	52847
X04325	31646
X044C7	29934
X04408 X04412	31914 35447
X04470	28638
X04503	36490
X04526	31667
X04571	31120
X04697	31991
X04707	31206
X05276	37201
X05332	35740
X05345	32457
X05606	28930
X05908 X06272	34387 30865
X06537	34234
X06617	36143
X06747	36101
X06834	5941
X07270	32435
X07429	29524
X07465	31769
X07695	34072
X07767	35478
X07819	35798
X07820	36628 33970
X07876 X12433	32451
X12451	29714
X12597	32326
X12654	35908
X12791	36112
X13238	1200056
X13425	31590
X13482	37546

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Table 6

ACC. NO.	GI NUMBER
X13585	37579
X13839	28329
X14420	30057
X14608	296365
X15183	32487
X15187	37260
X15653	37598
X15722	31824
X15729	38317
X15759	12583
X15822	30146
X15880	30029
X15949	33966
X16896	33800
X16940	36502
X17206	34391
X17620	35067
X51602	31431
X51668	6641
X52022	3127925
X52174	28321
X52235	34372
X52268	44100
X52882	311380
X52943	28912
X53280	29504
X53605	<b>35</b> 683
X53777	34198
X53793	28383
X53961	34415
X54199	31641
X54326	31957
X54473	311347
X54941	29976
X55525	30101
X55654	14015
X56351	28582
X56415	28410
X56807	30507
X56932	23690
X56958	4803677
X56998	37564
X56999	37568
X57025	33007
X57351	311373
X57352	311374
X57398	35526
X57500	35037
X57766	456256
X57810	33717
X57812	33723
X58141	28381
X59131	3776087
X59352	37214
X59405	34508
M59406	34509
X59618	36154

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Table 6

ACC. NO.	GI NUMBER
X59834	3183C
X59841	35314
X59892	3082C
X6C036	38261
M60111	34768
X6C221	509290
X6C489	31099
X6C656	31134
X60708	35335
X61100	38078
X61970	296739
X62048	499071
X62534	32332
X62744	
	36062
X63380	36168
X63422	12585
X63423	12587
X63432	28335
X63563	36121
X63564	36123
X63679	37264
X63753	36545
X64330	28934
X64707	29382
X64838	35998
X64875	398163
X65550	415818
X65644	38259
X65923	31302
X66276	36500
X66534	31683
X66785	30489
X67098	475908
X67698	37476
X67951	287640
X68277	29980
X68560	38417
X68684	296460
X69086	34811
X69111	32294
K69117	312394
X69711	296534
X70326	
X70394	38434
	468707
X71087	288396
X71129	297901
X72727	460788
X73114	402618
X73608	793844
X73874	439271
X73902	452754
X74070	395086
X74331	510 <b>4</b> 07
X74801	671526
X75304	405714
X75535	1644300
X75546	453156

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Table 6

ACC. NO. X75593 X75861 X76013 X76057 X76105 X76180 X76302 X76538 X76578 X76770 X76771 X77196 X77548 X77753 X78136 X78137 X78520 X78627 X78926 X78933 X79201 X79389 X79535 X79888 X80676 X80695 X80754 X80822 X80910 X80916 X81109 X81198 X81695 X81789 X81836 X81899 X81836 X81899 X81836 X81899 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X81989 X819	GI NUMBER 452319 456258 531595 416016 434844 452649 431952 434976 435144 2706486 556782 1905802 704462 469145 1524102 460772 460770 854101 607129 498724 498720 531105 510904 496886 780240 1240060 619490 577778 527577 531475 531367 535057 773574 940515 551449 577052 1702923 2274973 1310670 758065 1617117 1089849 639692 1502344 728715 854125 755749 1262344 806563 806565 899293
X85134 X85337 X85372	1262344 806563

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Table 6

ACC. NO.	GT 177
ACC. NO.	GI NUMBER
X39652	902619
X89673	902329
X89852	1050424
X90444	1066232
X90568	1212991
M90583	1071680
X90870	
	1103676
X91257	1050526
X91484	987789
X92098	1212964
X92689	1296629
X92845	2344811
X93972	1945270
X93036	1085025
M94323	1213612
X94754	1702931
X94910	3413292
X95240	
	1262818
X95325	1167837
M95384	2792003
X95677	1491701
X97065	
	1296665
X97324	1806039
X97335	1507823
X97674	1877214
X97675	
	1834512
X97676	1313923
X97677	1313925
X97772	1944613
X97790	1306089
∑97341	1322145
X97388	1666659
N98263	1770461
X98296	1666074
X98608	1419345
X98743	1498228
X98801	1419566
X98891	1420872
X38901	
	3928200
M99098	1592819
X99585	1770518
X99920	1694827
Y93052	
	30308
Y00281	36052
Y00282	36043
100345	35569
100361	37575
Y00503	34038
Y00711	34328
Y00815	34266
100971	35699
Y07593	1881446
Y07968	1552327
Y08319	1922312
Y03915	1877201
Y03991	1817583
Y09188	2230868

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## Table 6

ACC. NO.	GI NUMBER
Y09267	1834492
Y09305	1669684
Y09615	1707506
Y09836	1743253
Y10313	2706510
Y10351	3021391
Y11164	2145440
Y11395	2894085
Y11435	2910996
Y12065	2230877
Y12781	3021408
Y12860	2808530
Y13286	2853173
Y13323	5042231
Y13834	5327058 2315201
Y13936 Y14738	2765426
Y15286	2584788
Y15906	5327056
Y16132	2765526
Y16241	3378195
Y16414	2924334
Y16610	3273088
Y18314	5457356
211227	34926
211241	34939
Z11338	36236
211491	41072
211531	31103
211692	31107 36425
Z11793 Z13009	31072
214244	30150
214392	6057
214468	6133
215085	20243
215124	7226
Z15132	65134
Z15187 `	288856
Z16412	48235
216695	23264
Z17335	10476
217710	16894 30786
Z17864 Z18356	30768
Z18538	28711
Z19002	38517
Z19471	29313
Z20613	27358
Z21507	38521
222534	402184
Z22548	438068
Z22658	297411
222968	312141
723064	3256006
Z24611	394980
Z24724	505034

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## MISSING AT THE TIME OF PUBLICATION

Table 1-1

Clanal	م ما	4		100101111				
Cione				MPM-116/				Tumor Aver
788180	Tumors	BPH		Aver. BPH		Aver. BPH		
308682	25.18 8.99	0.50	102.19	5.44	15.38	90.33	40.12	50.69
646037		0.27	26.49	39.49	22.07	29.80	46.70	32.91
	24.42	0.75	53.07	14.65	28.67	17.92	49.22	32.70
281003	128.77	4.90	33.20	39.14	7.16	31.28	20.52	26.26
1034473	33.61	1.35	44.07	2.71	18.38	21.16	38.51	24.97
284701	33.05	1.53	17.19	22.80	11.69	43.48	13.22	21.68
66336	1 1	0.10	26.22	29.17	26.08	1.00	1.00	16.70
281039		0.90	1.80		1.52	66.03	1.32	15.57
26736		0.62	18.76	4.16	3.08	32.76	18.11	15.37
784168		3.83	4.81	12.77	16.92	15.79	18.91	13.84
486035		1.86	10.82	21.22	9.17	10.98	14.41	13.32
593185		2.29	3.25	8.22	9.64	26.71	13.34	12.23
222157		0.73	19.15	11.68	10.10		15.47	12.14
35058		1.44	4.36		6.63		6.94	10.81
730739		3.45	9.24		21.62		6.29	10.57
1636908		0.63	7.98		11.24		5.88	10.44
292515		1.56	8.49		6.85		12.51	10.38
450049		1.80	8.59		7.59		23,45	10.21
666334		0.16	9.09		10.25			10.14
32160		2.77	8.40		11.15			9.69
26519		3.76	10.15		8.76			9.59
42415	•		3.28		11.07			9.49
50772	l l		4.58		7.33			9.47
1412481			0.84					9.31
197525					7.12			9.01
490178			5.26					8.99
461522								8.74
746152	1							1
712950	1		L					8.58
814053	· ·							8.38
161044								a .
8258 <i>5</i> ° 59529°								
								t .
59569:			B:					<b>1</b>
48896		I .						1
25937- 28208								1
82394								
82394 44771								
104881								
4616								
46027								
68729								
75739 73129								
		•	1					
155652								
82610								
82573								
2216	20.2	4 2.8	0 5.5	54 5.0	5 9.7	71 9.2	0 6.7	7.24

Table 1-1

	!							
283173	17.85	2.47	17.57	3.98	4.39	7.32	2.81	7.21
825822	14.07	1.97	5.87	8.14	6.53	10.54	4.71	7.16
280217	20.52	2.89	9.65	0.93	2.71	20.73	1.45	7.09
262053	10.75	1.52	5.67	9.07	B.71	8.34	3.60	7.08
66317	5.50	0.78	15.41	4.56	4.00	10.34	0.99	7.06
565235	35.83	5.15	3.19	8.34	3.79	17.38	2.08	6.96
1468260	4.53	0.65	13.06	1.36	7.45	4.03	8.89	6.96
826103	1.31	0.19	8.19	7.96	5.94	9.88	2.70	6.93
796366	34.94	5.05	10.57	5.34	12.35	3.13	3.18	6.92
877835	23.06	3.34	9.71	3.97	5.61	7.41	7.82	6.90
258300	18.15	2.65	4.31	7.13	13.32	1.26	8.27	6.86
814062	0.81	0.12	10.04	5.59	8.22	8.85	1.42	6.83
1034738	22.40	3.29	5.92	5.75	6.61	9.70	6.12	6.82
951313	6.84	1.01	2.78	4.19	15.48	7.44	3.86	6.75
253132	1.40	0.21	8.70	9.55	4.09	5.72	5.23	6.66
824510	1.38	0.21	8.56	9.50	7.92	5.67	1.59	6.65
826072	1.20	0.18	7.01	8.64	4.02	10.85	2.42	6.59
796754	0.66	0.10	6.20	5.19	5.58	9.09	6.47	6.5C
251806	2.94	0.46	6.74	6.71	5.58	9.16	3.74	6.39
840306	2.49	0.39	2.46	5.82	8.43	9.48	5.73	6.38
815163	1.20	0.19	6.07	6.53	9.54	7.44	2.20	6.36
838285	1.34	0.21	4.71	7.31	6.15	8.05	5.54	6.35
565863	3.93	0.63	13.52	9.05	2.58	3.19	3.01	6.27
757435	365.78	58.35	3.90	3.17	17.54	3.19	3.55	6.27
67401	1.47	0.23	1.08	1.27	26.14	1.90	0.96	6.27
796239	26.07	4.18	1.88	2.16	10.01	7.23	9.94	6.24
321706	7.80	1.25	8.36	1.45	14.65	1.76	4.95	6.23
564801	12.35	1.99	5.89	2.75	17.66	3.44	1.34	6.21
9507 <b>78</b> 884 <b>60</b> 6	3.00 33.46	0.49 5.43	12.47	7.19	4.14	3.62	3.50	6.18
454317		1	6.01	3.92	16.90	2.25	1.74	6.16
796646	20.56 6.21	3.34 1.01	3.52 5.13	4.38	7.94	12.41	2.55	6.16
840894	34.81	5.66	2.34	3.61 6.76	5.34 5.31	6.22 9.85	10.48	6.16
52933	3.59	0.58	5.96				6.47	6.15
5949 <b>9</b> 4	14.61	2.38	5.57	6.27 4.29	4.93 8.11	6.70 6.75	6.87	6.14
586780		0.31	11.12	7.73	4.39	3.68	5.98 3.68	6.14
868282		1.47	5.69	6.02	6.96	6.08	5.84	6.12 6.12
1535596		66.91	5.47	9.25	6.43	4.48	4.39	6.00
259973			6.61	6.24	5.06	6.62	5.42	5.99
839081		1 1	10.16	10.63	1.88	5.07	2.14	5.98
359661	1		4.04	9.90	3.64	7.90	4.38	5.97
509731			4.92	3.91	8.19	7.51	5.20	5.95
770066			1.49	2.31	5.43	18.27	2.17	5.94
151055			5.93	10.61	2.93	4.74	5.38	5.92
842927			2.96	5.54	10.03	6.26	4.59	5.88
824126			8.58	7.60	6.12	6.06	1.00	5.87
814662			5.53	5.17	7.24	9.88	1.37	5.84
105617			6.48	4.11	1.30	15.30	1.99	5.84
84094			11.87	1.67	12.92	1.88	0.84	5.83
81158			8.01	4.53	6.69	4.59	5.11	5.78
163728			2.86	2.47	10.87	9.32	3.03	5.71
	•	•						

Table 1-1

838807	1.90	0.33	6.10	5.85	4.20	7.62	4.77	5.71
77915	14.99	2.63	3.34	8.82	3.66	2.66	10.02	5.70
252314	0.69	0.12	8 <i>.</i> 50	5.50	6.04	3.55	4.72	5.66
814235	12.37	2.19	6.17	6.36	5.75	7.43	2.52	5.65
461476	0.78	0.14	7.32	7.96	5.15	5.06	2.71	5.64
754358	12.77	2.26	4.29	5.26	8.83	2.50	7.34	5.64
53022	1.34	0.24	3.38	6.17	9.26	3.91	5.41	5.63
859228	85.06	15.20	5.32	4.79	6.57	7.30	4.01	5.60
290370	0.62	0.11	4.96	3.44	7.59	8.28	3.41	5.54
81331	2.84	0.52	1.67	4.49	1.34	18.34	1.80	5.53
260116	0.88	0.16	6.00	5.67	4.91	5.69	5.33	5.52
416374	4.49	0.81	3.17	8.84	3.61	10.29	1.67	5.52
840606	15.85	2.88	3.74	4.47	5.11	9.38	4.79	5.50
51773	11.48	2.09	6.89	4.08	3.70	8.77	4.03	5.50
240694	4.53	0.83	5.08	8.53	3.38	3.73	6.64	5.47
1605407	9.00	1.65	4.85	3.69	10.03	6.25	2.42	5.45
139771	7.72	1.42	13.10	8.46	1.72	1.70	2.24	5.44
1630990	20.64	3.80	8.87	4.51	3.52	9.31	0.98	5.44
1390584	6.54	1.21	3.13	4.93	5.15	12.83	0.99	5.41
186918	3.04	0.56	2.90	5.49	4.54	11.24	2.73	5.38
462097	8.32	1.55	7.30	2.88	1.25	9.95	5.50	5.38
32381	2.18	0.41	1.75	2.88	9.49	3.19	9.57	5.38
491764	0.81	0.15	6.00	5.07	5.46	5.33	4.98	5.37
1456160	41.61	7.76	2.23	3.67	7.39	11.23	2.29	5.36
159608	14.43	2.70	10.47	1.64	1.63	7.83	5.21	5.36
80050	8.15	1.52	6.41	4.16	11.93	0.95	3.31	5.35
610124	1.70	0.32	4.17	6.14	5.42	5.63	5.22	5.32
280763	0.53	0.10	3.60	5.23	5.40	6.52	5.64	5.28
283312	63.94	12.14	6.26	3.00	2.86	6.54	7.69	5.27
359781	7.82	1.49	3.10	9.11	3.52	6.46	4.12	5.26
297439	48.60	9.26	5.11	9.34	4.69	2.38	4.73	5.25
731410	, ,	0.10	5.74	4.50	5.23	5.26	5.48	5.24
772416		0.20	4.21	4.42	4.71	7.57	5.02	5.19
267419		0.19	3.98	9.05	3.92	5.50	3.38	5.16
897807		0.99	5.07	5.96	9.08	3.03	2.65	5.16
593658		0.22	4.01	5.03	5.25	5.65	5.29	5.04
511952		45.17	0.81	4.98	1.25	15.74	2.20	5.00
743452		1.59	7.90	7.46	2.03	4.53	2.98	4.98
878281		1.37	5.03	1.94	7.14	8.50	2.27	4.98
825297		0.19	4.48	3.92	7.31	4.87	4.28	4.97
731016	0.85	0.17	6.76	6.19	3.31	4.49	4.06	4.96
824723		2.84	3.64	3.92	5.23	8.15	3.87	4.96
824123		0.94	6.80	5.17	5.77	4.87	2.17	4.95
75523		3.62	5.25	3.08	3.43	8.49	4.50	4.95
78413		3.50	3.20	5.09	7.85	4.99	3.50	4.93
158554	1		5.67	6.28	3.85	4.93	3.89	4.92
70167	I .		7.12	3.55	3.14	8.18	2.54	4.91
81401			6.83	7.64	4.06	2.12	3.86	4.90
53018		1.74	2.68	5.31	6.79	3.57	6.04	4.88
77328		3.27	4.28	4.12	3.43	7.01	5.50	4.87
19472	•	1.07	8.66	7.32	2.79	2.88	2.66	4.86

Table 1-1

238661         2.74         0.56         9.41         7.84         2.13         2.69         2.21         4.8           773248         21.90         4.51         1.37         2.18         6.84         6.57         7.31         4.8           124252         6.06         1.25         11.29         6.83         1.86         2.10         2.21         4.8           814306         18.28         3.77         3.29         5.66         3.89         7.34         4.08         4.8           754280         23.83         4.92         9.79         5.07         5.33         1.97         2.08         4.8           461592         11.87         2.45         8.03         6.34         4.05         3.94         1.87         4.8           401592         11.87         2.45         8.03         6.34         4.05         3.94         1.81         4.8         4.09         4.8         4.8         4.05         3.94         1.8         4.8         4.3         7.93         4.8         4.8         4.3         3.93         1.8         4.93         1.1         8.0         3.94         1.8         4.9         4.1         4.9         4.1         4.9 <td< th=""><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th></td<>									
773248 2190	788655	7.66	1.58	4.19	3.33	7.19	6.38	3.22	4.86
124252	238661	2.74	0.56	9.41	7.84	2.13	2.69	2.21	4.86
814306         18.28         3.77         3.29         5.66         3.89         7.34         4.08         4.8           754280         23.83         4.92         9.79         5.07         5.33         1.97         2.08         4.8           1636108         3.58         0.74         5.59         4.80         5.19         4.56         4.09         4.8           461592         11.87         2.45         8.03         6.34         4.05         3.94         1.87         4.8           703943         4.77         0.99         3.01         6.34         6.79         4.06         3.91         4.8           289071         37.48         7.78         3.68         4.93         11.18         2.30         2.00         4.8           510576         132.22         27.52         5.44         2.70         2.95         8.26         4.68         4.3           511428         24.31         5.08         5.64         3.21         4.36         5.17         5.56         4.3           488431         10.48         2.19         6.83         4.34         3.39         2.92         6.45           823615         9.40         1.97         6.3		21.90	4.51	1.37	2.18	6.84	6.57	7.31	4.86
754280         23 83         4.92         9.79         5.07         5.33         1.97         2.08         4.8           1636108         3.58         0.74         5.59         4.80         5.19         4.56         4.09         4.8           461592         11.87         2.45         8.03         6.34         4.05         3.94         1.87         4.8           771053         16.14         3.34         3.85         4.29         4.79         3.56         7.67         4.8           703943         4.77         0.99         3.01         6.34         6.79         4.06         3.91         4.8           823691         7.42         1.54         2.71         4.70         4.41         8.34         3.93         4.8           510576         132.22         27.52         5.44         2.70         2.95         8.26         4.68         4.1           510576         132.22         27.52         5.44         2.70         2.95         8.26         4.68         4.1           488431         10.48         2.19         6.83         4.31         3.39         2.92         6.45         4.1           4882311         10.48         2.									4.85
1636108   3.58   0.74   5.59   4.80   5.19   4.56   4.00   4.8							7.34		4.85
461592 11.87 2.45 8.03 6.34 4.05 3.94 1.87 4.8 771053 16.14 3.34 3.85 4.29 4.79 3.56 7.67 7.67 7.03943 4.77 0.99 3.01 6.34 6.79 4.06 3.91 4.8 7.03943 4.77 0.99 3.01 6.34 6.79 4.06 3.91 4.8 7.03943 4.77 0.99 3.01 6.34 6.79 4.06 3.91 4.8 7.03943 4.77 8.5 7.78 3.68 4.93 11.18 2.30 2.00 4.8 7.03943 7.78 5.05 7.0 1.05 5.25 5.44 2.70 2.95 8.26 4.68 4.3 5.05 7.0 1.05 5.25 4.74 5.40 3.85 4.73 5.11428 24.31 5.08 5.64 3.21 4.36 5.17 5.56 4.5 888431 10.48 2.19 6.83 4.34 3.39 2.92 6.45 4.3 359701 5.21 1.09 4.50 5.09 3.54 3.24 7.52 4.5 1606837 23.76 4.98 6.95 2.67 3.34 5.80 5.10 4.8 23615 9.40 1.97 6.36 2.61 10.35 2.83 1.67 4.8 23615 9.40 1.97 6.36 2.61 10.35 2.83 1.67 4.8 13275 1.75 0.37 2.37 3.77 3.58 8.89 5.04 4.8 13275 1.75 0.37 2.37 3.77 3.58 8.89 5.04 4.8 375682 31.04 6.60 3.43 4.82 7.07 3.67 4.53 4.8 23615 0.73 1.08 1.71 5.35 8.20 6.29 2.02 3.75682 31.04 6.60 3.43 4.82 7.07 3.67 4.53 4.8 23455 0.73 0.16 4.68 5.34 5.04 4.18 4.15 4.27 0.74 0.16 4.49 4.65 4.35 5.04 4.18 4.15 4.27 0.74 0.16 4.49 4.65 4.35 5.04 4.18 4.15 4.27 0.74 0.16 4.49 4.65 4.35 5.04 4.18 4.15 4.15 4.29 0.92 0.87 0.19 5.08 4.85 5.04 4.18 4.15 4.15 4.29 0.92 0.87 0.19 5.08 4.85 5.04 4.18 4.15 4.15 4.29 0.92 0.87 0.19 5.08 4.85 5.04 4.2 3.5 4.49 6.81 1.39 0.93 0.25 6.55 4.44 2.37 4.47 9.42 2.38 8.4878 2.58 1 5.59 2.71 4.00 3.35 9.63 3.38 4.20 0.94 1.18 4.15 4.15 4.29 0.92 0.87 0.19 5.08 4.85 3.34 5.04 4.18 4.15 4.15 4.29 0.26 3.47 5.67 4.78 4.79 4.2 2.38 8.4878 2.58 1 5.59 2.71 4.00 3.35 9.63 3.38 4.30 0.10 1.17 4.29 3.24 4.06 5.34 4.18 4.15 4.15 4.29 0.26 3.47 5.67 4.78 4.79 4.2 2.38 4.48 1.16 6.06 6.81 1.21 0.26 6.55 4.44 2.37 4.47 9.42 2.38 8.4878 2.58 1 5.59 2.71 4.00 3.35 9.63 3.38 4.39 0.29 1.38 1.38 1.39 5.00 1.10 1.74 7.29 3.24 7.49 3.02 1.39 2.60 5.81 1.39 1.024 6.19 2.21 2.17 2.11 2.11 6.05 6.55 1.44 2.37 4.47 9.42 2.38 4.40 5.31 1.38 1.39 5.00 1.10 1.74 7.29 3.24 7.49 3.02 1.39 2.50 1.24 4.62 2.07 5.28 2.80 7.76 1.30 2.50 6.55 1.44 4.2 2.37 4.47 9.42 2.38 4.40 1.39 1.29 1.245 5.25 1.18 3.96 4.49 6.00 3.91 3.98 5.73 1.155 4.29 0.55 1.18 3.96		23.83	4.92	9.79		5.33			4.85
771053	,	3.58			4.80	5.19			4.85
703943					6.34				4.84
823691         7.42         1.54         2.71         4.70         4.41         8.34         3.93         4.8           289071         37.78         7.78         3.68         4.93         11.18         2.30         2.00         4.8           510576         132.22         27.52         5.44         2.70         2.95         8.26         4.68         4.3           264465         0.77         0.16         5.25         4.74         5.40         3.85         4.73         4.7           511428         24,31         5.08         5.64         3.21         4.36         5.17         5.56         4.2           488431         10.48         2.19         4.50         5.09         3.54         3.24         7.52         4.7           359701         5.21         1.09         4.50         5.09         3.54         3.24         7.52         4.7           323615         9.40         1.97         6.36         2.61         10.35         2.83         1.67         4.2           795265         4.58         0.97         1.65         3.44         3.71         12.61         2.30         4.1         4.1         4.9         4.6         3.6									4.83
289071   37.48   7.78   3.68   4.93   11.18   2.30   2.00   4.8									4.82
510576         132.22         27.52         5.44         2.70         2.95         8.26         4.68         4.3           264465         0.77         0.16         5.25         4.74         5.40         3.85         4.73         4.7           511428         24.31         5.08         5.64         3.21         4.36         5.17         5.56         4.7           488431         10.48         2.19         6.83         4.34         3.39         2.92         6.45         4.7           359701         5.21         1.09         4.50         5.09         3.54         3.24         7.52         4.1           1606837         23.76         4.98         6.95         2.67         3.34         5.80         5.10         4.2           823615         9.40         1.97         6.36         2.61         10.35         2.83         1.67         4.           795265         4.58         0.97         1.65         3.44         3.71         12.61         2.30         4.           796351         2.34         0.50         5.27         6.47         3.71         4.50         3.66         4.           787938         5.09         1.08									4.82
264465									4.82
511428         24.31         5.08         5.64         3.21         4.36         5.17         5.56         4.5           488431         10.48         2.19         6.83         4.34         3.39         2.92         6.45         4.5           359701         5.21         1.09         4.50         5.09         3.54         3.24         7.52         4.5           1606837         23.76         4.98         6.95         2.67         3.34         5.80         5.10         4.           823615         9.40         1.97         6.36         2.61         10.35         2.83         1.67         4.           795265         4.58         0.97         1.65         3.44         3.71         12.61         2.30         4.           796351         2.34         0.50         5.27         6.47         3.71         4.50         3.66         4.           787938         5.09         1.08         1.71         5.35         8.20         6.29         2.02         4.           375682         31.04         6.60         3.43         4.82         7.07         3.67         4.53         4.         2.54         4.         2.54         4.									4.81
488431         10.48         2.19         6.83         4.34         3.39         2.92         6.45         4.5           339701         5.21         1.09         4.50         5.09         3.54         3.24         7.52         4.5           1606837         23.76         4.98         6.95         2.61         10.35         2.83         1.67         4.6           823615         9.40         1.97         6.36         2.61         10.35         2.83         1.67         4.6           795265         4.58         0.97         1.65         3.44         3.71         12.61         2.30         4.           796351         2.34         0.50         5.27         6.47         3.71         4.50         3.64         4.           787938         5.09         1.08         1.71         5.35         8.20         6.29         2.02         4.           375682         31.04         6.60         3.43         4.82         7.07         3.67         4.53         4.           869450         51.58         10.98         10.12         2.49         0.92         4.42         5.54         4.         1.         5.54         4.         1.									4.79
359701   5.21   1.09   4.50   5.09   3.54   3.24   7.52   4.50   1.00   4.50   5.09   3.54   3.24   7.52   4.50   1.00   4.50   5.09   3.54   3.24   7.52   4.50   4.50   5.00   4.50   5.00   4.50   5.00   4.50   5.00   4.50   5.00   4.50   5.00   4.50   5.00   4.50   5.00   4.50   5.00   4.50   5.00   4.50   5.00   4.50   5.00   4.50   5.00   4.50   5.00   4.50   5.00   5.27   6.47   3.71   12.61   2.30   4.50   5.00   5.27   6.47   3.71   4.50   3.66   4.50   5.00   5.27   6.47   3.71   4.50   3.66   4.50   5.50   5.27   6.47   3.71   4.50   3.66   4.50   3.75   5.50   5.27   6.47   3.71   4.50   3.66   4.50   3.75   5.50   5.27   6.47   3.71   4.50   3.66   4.50   3.75   5.50   4.50   5.50   5.27   6.47   3.71   4.50   3.66   4.50   3.75   5.50   4.50   5.50   5.27   6.47   3.71   4.50   3.66   4.50   3.75   5.50   4.50   5.50   4.50   5.50   4.50   5.50   4.50   5.50   4.50   5.50   4.50   5.50   4.50   5.50   4.50   5.50   4.50   5.50   4.50   5.50   4.50   5.50   4.50   5.50   4.50   5.50   4.50   5.50   4.50   5.50   4.50   5.50   4.50   5.50   4.50   5.50   4.50   5.50   4.50   5.50   4.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50   5.50			4					ľ	4.79
1606837   23.76   4.98   6.95   2.67   3.34   5.80   5.10   4.232615   9.40   1.97   6.36   2.61   10.35   2.83   1.67   4.230   4.813275   1.75   0.37   2.37   3.77   3.58   8.89   5.04   4.796351   2.34   0.50   5.27   6.47   3.71   4.50   3.66   4.796351   2.34   0.50   5.27   6.47   3.71   4.50   3.66   4.796351   3.104   6.60   3.43   4.82   7.07   3.67   4.53   4.869450   51.58   10.98   10.12   2.49   0.92   4.42   5.54   4.82   2.51435   0.73   0.16   4.68   5.34   5.04   4.18   4.15   4.82   4.207   0.74   0.16   4.49   4.65   4.35   4.86   5.03   4.82   2.51517   0.73   0.16   4.87   4.91   5.29   3.94   4.18   4.15   4.66631   1.21   0.26   3.47   5.67   4.78   4.73   4.50   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.35   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86   5.34   4.86									4.78
823615         9.40         1.97         6.36         2.61         10.35         2.83         1.67         4.           795265         4.58         0.97         1.65         3.44         3.71         12.61         2.30         4.           813275         1.75         0.37         3.73         3.58         8.89         5.04         4.           796351         2.34         0.50         5.27         6.47         3.71         4.50         3.66         4.           787938         5.09         1.08         1.71         5.35         8.20         6.29         2.02         4.           375682         31.04         6.60         3.43         4.82         7.07         3.67         4.53         4.53         4.53         4.53         4.53         4.53         4.22         5.54         4.         4.53         4.86         5.03         4.         4.18         4.15         4.         4.15         4.         4.52         4.42         5.54         4.         4.18         4.15         4.         4.         4.53         4.86         5.03         4.         4.         4.65         4.35         4.86         5.03         4.         4.         4.65									4.78
795265         4.58         0.97         1.65         3.44         3.71         12.61         2.30         4.           813275         1.75         0.37         2.37         3.77         3.58         8.89         5.04         4.           796351         2.34         0.50         5.27         6.47         3.71         4.50         3.66         4.           787938         5.09         1.08         1.71         5.35         8.20         6.29         2.02         4.           375682         31.04         6.60         3.43         4.82         7.07         3.67         4.53         4.68         6.99         2.02         4.42         5.54         4.         4.53         4.68         5.34         5.04         4.18         4.15         4.         4.55         5.44         4.55         5.54         4.         4.55         5.54         4.         4.55         5.54         4.         4.55         5.54         4.         4.55         5.54         4.         4.55         5.54         4.         4.55         5.34         5.04         4.18         4.         4.55         5.34         4.         4.55         3.84         4.56         5.33         4. <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td>4.77</td>									4.77
813275         1.75         0.37         2.37         3.77         3.58         8.89         5.04         4.78         4.79         4.50         3.66         4.78         4.79         4.50         3.66         4.78         4.50         3.66         4.78         4.51         4.50         3.66         4.42         5.56         4.42         5.54         4.53         4.53         4.53         4.53         4.53         4.53         4.53         4.53         4.53         4.53         4.53         4.53         4.53         4.53         4.53         4.53         4.53         4.53         4.42         5.54         4.53         4.65         5.34         5.04         4.18         4.15         4.42         5.54         4.42         5.54         4.42         5.54         4.42         5.54         4.42         5.54         4.42         5.54         4.42         5.54         4.42         5.54         4.42         5.54         4.42         5.54         4.42         5.54         4.42         5.54         4.42         5.54         4.42         5.54         4.42         5.54         4.42         5.54         4.42         5.54         4.42         5.54         4.42         5.54         4.42         <									4.76
796351         2.34         0.50         5.27         6.47         3.71         4.50         3.66         4.787938         5.09         1.08         1.71         5.35         8.20         6.29         2.02         4.8877682         31.04         6.60         3.43         4.82         7.07         3.67         4.53         4.869450         51.58         10.98         10.12         2.49         0.92         4.42         5.54         4.54         4.55         4.86         5.54         4.62         5.54         4.62         5.54         4.62         5.54         4.62         5.54         4.62         4.63         5.34         5.04         4.18         4.15         4.42         5.54         4.62         4.35         4.86         5.03         4.84         4.65         4.35         4.86         5.03         4.86         5.03         4.87         4.91         5.29         3.94         4.18         4.66         5.03         4.88         4.91         5.29         3.94         4.18         4.66         5.03         4.88         4.91         5.29         3.94         4.18         4.66         5.03         4.48         4.91         5.29         3.94         4.18         4.49         4.06         5.									4.74
787938         5.09         1.08         1,71         5.35         8.20         6.29         2.02         4.375682         31.04         6.60         3.43         4.82         7.07         3.67         4.53         4.53         4.53         869450         51.58         10.98         10.12         2.49         0.92         4.42         5.54         4         4.25         4.51         4         4.25         5.54         4         4.53         4.18         4.15         4         4.22         4.67         4.01         4.68         5.34         5.04         4.18         4.15         4         4.22         4.67         4.01         6.01         4.49         4.65         4.35         4.86         5.03         4         4.18         4.15         4         4.65         4.35         4.86         5.03         4         4.21         7.01         6.02         7.01         6.02         7.01         6.02         7.01         6.02         7.01         6.02         7.02         7.02         7.02         8.26         6.21         3.21         7.02         8.29         3.94         4.18         4.42         7.02         4.47         7.03         4.47         7.03         7.02         7.02									4.73
375682         31.04         6.60         3.43         4.82         7.07         3.67         4.53         4.8869450         51.58         10.98         10.12         2.49         0.92         4.42         5.54         4.62         251435         0.73         0.16         4.68         5.34         5.04         4.18         4.15         4.82         4.67         1.00         4.49         4.65         4.35         4.86         5.03         4         4.18         4.15         4.86         5.03         4         4.18         4.15         4.86         5.03         4         4.18         4.15         4.86         5.03         4         4.18         4.15         4.86         5.03         4         4.18         4.18         4.66         6.00         6.10         1.00         9.34         7.13         2.17         3.26         1.38         4         4.66         6.93         4.85         3.84         4.06         5.34         4         4.86         6.93         4.85         3.84         4.06         5.34         4         4.66         6.11         1.32         8.25         6.21         3.19         2.69         2.81         4.50         4.73         4.50         4.73         4.50								1	4.72
869450         51.58         10.98         10.12         2.49         0.92         4.42         5.54         4           251435         0.73         0.16         4.68         5.34         5.04         4.18         4.15         4           824267         0.74         0.16         4.49         4.65         4.35         4.86         5.03         4           214006         4.67         1.00         9.34         7.13         2.17         3.26         1.38         4           251517         0.73         0.16         4.87         4.91         5.29         3.94         4.18         4           609052         0.87         0.19         5.08         4.85         3.84         4.06         5.34         4           1466631         1.21         0.26         3.47         5.67         4.78         4.73         4.50         4           1500000         6.11         1.32         8.25         6.21         3.19         2.69         2.81         4           1055201         30.25         6.55         4.44         2.37         4.47         9.42         2.38         4           840878         25.81         5.59         2.7									4.71
251435         0.73         0.16         4.68         5.34         5.04         4.18         4.15         4           824267         0.74         0.16         4.49         4.65         4.35         4.86         5.03         4           214006         4.67         1.00         9.34         7.13         2.17         3.26         1.38         4           251517         0.73         0.16         4.87         4.91         5.29         3.94         4.18         4           609052         0.87         0.19         5.08         4.85         3.84         4.06         5.34         4           1466631         1.21         0.26         3.47         5.67         4.78         4.73         4.50           1500000         6.11         1.32         8.25         6.21         3.19         2.69         2.81           1055201         30.25         6.55         4.44         2.37         4.47         9.42         2.38           840878         25.81         5.59         2.71         4.00         3.35         9.63         3.38           773204         21.34         4.63         2.70         3.84         5.75         1.24									4.70
824267         0.74         0.16         4.49         4.65         4.35         4.86         5.03         4           214006         4.67         1.00         9.34         7.13         2.17         3.26         1.38         4           251517         0.73         0.16         4.87         4.91         5.29         3.94         4.18         4           609052         0.87         0.19         5.08         4.85         3.84         4.06         5.34         4           1466631         1.21         0.26         3.47         5.67         4.78         4.73         4.50         4           1500000         6.11         1.32         8.25         6.21         3.19         2.69         2.81         4           1055201         30.25         6.55         4.44         2.37         4.47         9.42         2.38         4           840878         25.81         5.59         2.71         4.00         3.35         9.63         3.38         4           773204         21.34         4.63         2.70         3.84         5.75         1.24         9.53         2.13         4         134969         6.81         1.49         10									4.70
214006       4.67       1.00       9.34       7.13       2.17       3.26       1.38       4         251517       0.73       0.16       4.87       4.91       5.29       3.94       4.18       4         609052       0.87       0.19       5.08       4.85       3.84       4.06       5.34       4         1466631       1.21       0.26       3.47       5.67       4.78       4.73       4.50       4         1500000       6.11       1.32       8.25       6.21       3.19       2.69       2.81       4         1055201       30.25       6.55       4.44       2.37       4.47       9.42       2.38       4         840878       25.81       5.59       2.71       4.00       3.35       9.63       3.38       7         773204       21.34       4.63       2.70       3.84       5.75       1.24       9.53       4         290841       6.35       1.38       7.54       6.98       2.99       3.39       2.13       4         824376       0.49       0.11       5.18       4.11       6.01       6.67       0.94       4         26617       4.62 <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td>4.68</td>									4.68
251517         0.73         0.16         4.87         4.91         5.29         3.94         4.18         4           609052         0.87         0.19         5.08         4.85         3.84         4.06         5.34         4           1466631         1.21         0.26         3.47         5.67         4.78         4.73         4.50         4           1500000         6.11         1.32         8.25         6.21         3.19         2.69         2.81         4           1055201         30.25         6.55         4.44         2.37         4.47         9.42         2.38         4           840878         25.81         5.59         2.71         4.00         3.35         9.63         3.38         4           773204         21.34         4.63         2.70         3.84         5.75         1.24         9.53         2           290841         6.35         1.38         7.54         6.98         2.99         3.39         2.13         4           824376         0.49         0.11         5.18         4.11         6.01         6.67         0.94         4           768643         41.57         9.10         2.42									4.68 4.66
609052         0.87         0.19         5.08         4.85         3.84         4.06         5.34         4           1466631         1.21         0.26         3.47         5.67         4.78         4.73         4.50         4           1500000         6.11         1.32         8.25         6.21         3.19         2.69         2.81         4           1055201         30.25         6.55         4.44         2.37         4.47         9.42         2.38         4           840878         25.81         5.59         2.71         4.00         3.35         9.63         3.38         4           773204         21.34         4.63         2.70         3.84         5.75         1.24         9.53         4           290841         6.35         1.38         7.54         6.98         2.99         3.39         2.13         4           824376         0.49         0.11         5.18         4.11         6.01         6.67         0.94         4           768643         41.57         9.10         2.42         4.02         11.03         2.91         2.45         4           26617         4.62         1.01         1.74									4.64
1466631         1.21         0.26         3.47         5.67         4.78         4.73         4.50         4           1500000         6.11         1.32         8.25         6.21         3.19         2.69         2.81         4           1055201         30.25         6.55         4.44         2.37         4.47         9.42         2.38         4           840878         25.81         5.59         2.71         4.00         3.35         9.63         3.38         4           773204         21.34         4.63         2.70         3.84         5.75         1.24         9.53         4           290841         6.35         1.38         7.54         6.98         2.99         3.39         2.13         4           134969         6.81         1.49         10.24         6.19         2.21         2.17         2.11         4           824376         0.49         0.11         5.18         4.11         6.01         6.67         0.94         4           768643         41.57         9.10         2.42         4.02         11.03         2.91         2.45         4         2.45         4         4         4         6.10									4.63
1500000         6.11         1.32         8.25         6.21         3.19         2.69         2.81         4           1055201         30.25         6.55         4.44         2.37         4.47         9.42         2.38         4           840878         25.81         5.59         2.71         4.00         3.35         9.63         3.38         4           773204         21.34         4.63         2.70         3.84         5.75         1.24         9.53         4           290841         6.35         1.38         7.54         6.98         2.99         3.39         2.13         4           134969         6.81         1.49         10.24         6.19         2.21         2.17         2.11         4           824376         0.49         0.11         5.18         4.11         6.01         6.67         0.94         4           768643         41.57         9.10         2.42         4.02         11.03         2.91         2.45         4           26617         4.62         1.01         1.74         7.29         3.24         7.49         3.02         3         30538         11.14         4.62         2.07         5									4.63
1055201         30.25         6.55         4.44         2.37         4.47         9.42         2.38         4           840878         25.81         5.59         2.71         4.00         3.35         9.63         3.38         4           773204         21.34         4.63         2.70         3.84         5.75         1.24         9.53         4           290841         6.35         1.38         7.54         6.98         2.99         3.39         2.13         4           134969         6.81         1.49         10.24         6.19         2.21         2.17         2.11         4           824376         0.49         0.11         5.18         4.11         6.01         6.67         0.94         4           768643         41.57         9.10         2.42         4.02         11.03         2.91         2.45         4           26617         4.62         1.01         1.74         7.29         3.24         7.49         3.02         4         3         3.02         4         4         4         4         4         4         4         4         4         4         4         4         4         4         4		1 1							4.63
840878     25.81     5.59     2.71     4.00     3.35     9.63     3.38     4       773204     21.34     4.63     2.70     3.84     5.75     1.24     9.53     4       290841     6.35     1.38     7.54     6.98     2.99     3.39     2.13     4       134969     6.81     1.49     10.24     6.19     2.21     2.17     2.11     4       824376     0.49     0.11     5.18     4.11     6.01     6.67     0.94       768643     41.57     9.10     2.42     4.02     11.03     2.91     2.45       26617     4.62     1.01     1.74     7.29     3.24     7.49     3.02       305538     11.44     2.54     6.22     5.46     2.58     2.41     5.84       81162     6.36     1.41     4.62     2.07     5.28     2.80     7.76       1637343     5.03     1.12     3.26     4.54     4.14     6.66     3.91       197779     8.61     1.91     10.37     7.03     1.68     1.51     1.91       815549     2.30     0.51     6.84     5.83     3.95     4.46     1.38       51700     3.66									4.62
773204         21.34         4.63         2.70         3.84         5.75         1.24         9.53         4           290841         6.35         1.38         7.54         6.98         2.99         3.39         2.13         4           134969         6.81         1.49         10.24         6.19         2.21         2.17         2.11         4           824376         0.49         0.11         5.18         4.11         6.01         6.67         0.94         4           768643         41.57         9.10         2.42         4.02         11.03         2.91         2.45         4           26617         4.62         1.01         1.74         7.29         3.24         7.49         3.02         4           305538         11.44         2.54         6.22         5.46         2.58         2.41         5.84         5.84         5.84         81162         6.36         1.41         4.62         2.07         5.28         2.80         7.76         1637343         5.03         1.12         3.26         4.54         4.14         6.66         3.91         19779         8.61         1.91         10.37         7.03         1.68         1.51		1 1							4.62
290841     6.35     1.38     7.54     6.98     2.99     3.39     2.13       134969     6.81     1.49     10.24     6.19     2.21     2.17     2.11       824376     0.49     0.11     5.18     4.11     6.01     6.67     0.94       768643     41.57     9.10     2.42     4.02     11.03     2.91     2.45       26617     4.62     1.01     1.74     7.29     3.24     7.49     3.02       305538     11.44     2.54     6.22     5.46     2.58     2.41     5.84       811162     6.36     1.41     4.62     2.07     5.28     2.80     7.76       1637343     5.03     1.12     3.26     4.54     4.14     6.66     3.91       197779     8.61     1.91     10.37     7.03     1.68     1.51     1.91       815549     2.30     0.51     6.84     5.83     3.95     4.46     1.38       51700     3.66     0.82     4.47     4.07     4.99     3.08     5.73       757206     5.25     1.18     3.96     4.49     6.00     3.91     3.98       838761     2.59     0.58     6.93     6.35     2		1 1							4.61
134969     6.81     1.49     10.24     6.19     2.21     2.17     2.11     4       824376     0.49     0.11     5.18     4.11     6.01     6.67     0.94     4       768643     41.57     9.10     2.42     4.02     11.03     2.91     2.45     4       26617     4.62     1.01     1.74     7.29     3.24     7.49     3.02     4       305538     11.44     2.54     6.22     5.46     2.58     2.41     5.84       811162     6.36     1.41     4.62     2.07     5.28     2.80     7.76       1637343     5.03     1.12     3.26     4.54     4.14     6.66     3.91       197779     8.61     1.91     10.37     7.03     1.68     1.51     1.91       815549     2.30     0.51     6.84     5.83     3.95     4.46     1.38       51700     3.66     0.82     4.47     4.07     4.99     3.08     5.73       757206     5.25     1.18     3.96     4.49     6.00     3.91     3.98       838761     2.59     0.58     6.93     6.35     2.56     3.70     2.80       343987     11.55		1 1							4.61
824376         0.49         0.11         5.18         4.11         6.01         6.67         0.94         4           768643         41.57         9.10         2.42         4.02         11.03         2.91         2.45         4           26617         4.62         1.01         1.74         7.29         3.24         7.49         3.02         3.02           305538         11.44         2.54         6.22         5.46         2.58         2.41         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84         5.84									4.58
768643         41.57         9.10         2.42         4.02         11.03         2.91         2.45           26617         4.62         1.01         1.74         7.29         3.24         7.49         3.02           305538         11.44         2.54         6.22         5.46         2.58         2.41         5.84           811162         6.36         1.41         4.62         2.07         5.28         2.80         7.76           1637343         5.03         1.12         3.26         4.54         4.14         6.66         3.91           197779         8.61         1.91         10.37         7.03         1.68         1.51         1.91           815549         2.30         0.51         6.84         5.83         3.95         4.46         1.38           51700         3.66         0.82         4.47         4.07         4.99         3.08         5.73           757206         5.25         1.18         3.96         4.49         6.00         3.91         3.98           838761         2.59         0.58         6.93         6.35         2.56         3.70         2.80           343987         11.55         2.5		1 .						1	4.58
26617     4.62     1.01     1.74     7.29     3.24     7.49     3.02       305538     11.44     2.54     6.22     5.46     2.58     2.41     5.84       811162     6.36     1.41     4.62     2.07     5.28     2.80     7.76       1637343     5.03     1.12     3.26     4.54     4.14     6.66     3.91       197779     8.61     1.91     10.37     7.03     1.68     1.51     1.91       815549     2.30     0.51     6.84     5.83     3.95     4.46     1.38       51700     3.66     0.82     4.47     4.07     4.99     3.08     5.73       757206     5.25     1.18     3.96     4.49     6.00     3.91     3.98       838761     2.59     0.58     6.93     6.35     2.56     3.70     2.80       343987     11.55     2.59     2.47     4.92     6.23     2.51     6.15       767406     0.44     0.10     5.11     3.93     4.22     4.46     4.52			1					2.45	4.57
811162     6.36     1.41     4.62     2.07     5.28     2.80     7.76       1637343     5.03     1.12     3.26     4.54     4.14     6.66     3.91       197779     8.61     1.91     10.37     7.03     1.68     1.51     1.91       815549     2.30     0.51     6.84     5.83     3.95     4.46     1.38       51700     3.66     0.82     4.47     4.07     4.99     3.08     5.73       757206     5.25     1.18     3.96     4.49     6.00     3.91     3.98       838761     2.59     0.58     6.93     6.35     2.56     3.70     2.80       343987     11.55     2.59     2.47     4.92     6.23     2.51     6.15       767406     0.44     0.10     5.11     3.93     4.22     4.46     4.52	2661		1.01				7.49	3.02	4.56
1637343         5.03         1.12         3.26         4.54         4.14         6.66         3.91           197779         8.61         1.91         10.37         7.03         1.68         1.51         1.91           815549         2.30         0.51         6.84         5.83         3.95         4.46         1.38           51700         3.66         0.82         4.47         4.07         4.99         3.08         5.73           757206         5.25         1.18         3.96         4.49         6.00         3.91         3.98           838761         2.59         0.58         6.93         6.35         2.56         3.70         2.80           343987         11.55         2.59         2.47         4.92         6.23         2.51         6.15           767406         0.44         0.10         5.11         3.93         4.22         4.46         4.52	30553		2.54	6.22	5.46	2.58	2.41	5.84	4.50
197779         8.61         1.91         10.37         7.03         1.68         1.51         1.91           815549         2.30         0.51         6.84         5.83         3.95         4.46         1.38           51700         3.66         0.82         4.47         4.07         4.99         3.08         5.73           757206         5.25         1.18         3.96         4.49         6.00         3.91         3.98           838761         2.59         0.58         6.93         6.35         2.56         3.70         2.80           343987         11.55         2.59         2.47         4.92         6.23         2.51         6.15           767406         0.44         0.10         5.11         3.93         4.22         4.46         4.52	81116	6.36	1.41	4.62	2.07	5.28	2.80	7.76	4.50
815549     2.30     0.51     6.84     5.83     3.95     4.46     1.38       51700     3.66     0.82     4.47     4.07     4.99     3.08     5.73       757206     5.25     1.18     3.96     4.49     6.00     3.91     3.98       838761     2.59     0.58     6.93     6.35     2.56     3.70     2.80       343987     11.55     2.59     2.47     4.92     6.23     2.51     6.15       767406     0.44     0.10     5.11     3.93     4.22     4.46     4.52	163734	5.03	1.12	3.26	4.54	4.14	6.66		4.50
51700         3.66         0.82         4.47         4.07         4.99         3.08         5.73           757206         5.25         1.18         3.96         4.49         6.00         3.91         3.98           838761         2.59         0.58         6.93         6.35         2.56         3.70         2.80           343987         11.55         2.59         2.47         4.92         6.23         2.51         6.15           767406         0.44         0.10         5.11         3.93         4.22         4.46         4.52	19777	9 8.61	1.91	10.37	7.03	1.68	1.51		4.50
757206         5.25         1.18         3.96         4.49         6.00         3.91         3.98           838761         2.59         0.58         6.93         6.35         2.56         3.70         2.80           343987         11.55         2.59         2.47         4.92         6.23         2.51         6.15           767406         0.44         0.10         5.11         3.93         4.22         4.46         4.52									4.49
838761     2.59     0.58     6.93     6.35     2.56     3.70     2.80       343987     11.55     2.59     2.47     4.92     6.23     2.51     6.15       767406     0.44     0.10     5.11     3.93     4.22     4.46     4.52									4.47
343987 11.55 2.59 2.47 4.92 6.23 2.51 6.15 767406 0.44 0.10 5.11 3.93 4.22 4.46 4.52									4.47
767406 0.44 0.10 5.11 3.93 4.22 4.46 4.52									4.47
									4.46
700271 27 581 8 531 2 99 921 601 270 2 121									4.45
2.00 0.01 3.00 0.01 2.12	790	32 37.58	8.53	2.88	8.31	5.01	3.70	2.12	4.41

Table 1-1

815036         1.62         0.37         5.65         5.86         3.71         4.71         2.08         4.40           201282         6.42         1.46         1.85         6.04         4.99         6.43         2.63         4.39           773208         26.40         6.05         2.45         3.37         5.47         1.22         9.32         4.37           238821         6.54         1.50         5.59         1.57         4.76         0.92         8.90         4.35           814798         32.86         7.58         3.15         4.62         2.68         9.26         1.99         4.34           196650         10.93         2.52         4.17         6.60         2.20         6.71         1.99         4.33           197373         35.22         8.15         4.78         2.95         2.51         8.35         3.04         4.32           200599         13.42         3.11         5.88         2.44         8.89         1.51         2.87         4.32           487733         15.01         3.48         3.75         5.14         4.04         6.05         2.61         4.32           253411         3.22			_						
201282         6.42         1.46         1.85         6.04         4.99         6.43         2.63         4.39           773208         26.40         6.05         2.45         3.37         5.47         1.22         9.32         4.37           238821         6.54         1.50         5.59         1.57         4.76         0.92         8.90         4.35           196650         10.93         2.52         4.17         6.60         2.20         6.71         1.99         4.33           971367         35.22         8.15         4.78         2.95         2.51         8.35         3.04         4.32           200599         13.42         3.11         5.88         2.44         8.89         1.51         2.87         4.32           487733         15.01         3.48         3.75         5.14         4.04         6.05         2.61         4.32           564537         1.79         0.42         2.77         5.08         5.29         4.04         4.38         4.31           50606         14.94         3.47         5.20         2.99         5.04         3.92         4.99         4.31           253314         1.52         0					5.86	3.71	4.71	2.08	4.40
201282 6.42 1.46 1.85 6.04 4.99 6.43 2.63 4.39 1.71 1.73 1.22 9.32 4.37 1.73 1.24 1.50 6.05 2.45 3.37 5.47 1.22 9.32 4.37 1.21 1.21 1.25 6.34 1.50 5.59 1.57 4.76 0.92 8.90 4.35 1.47 1.22 1.23 2.26 1.20 1.25 1.25 1.27 4.76 0.92 8.90 4.35 1.47 1.22 1.25 1.25 1.25 1.25 1.25 1.25 1.25				4.08	4.25		4.82	4.45	4.40
773208   26.40   6.05   2.45   3.37   5.47   1.22   9.32   4.37   121436   6.37   1.46   7.91   6.01   2.52   3.24   2.09   4.36   1814798   32.86   7.58   3.15   4.62   2.68   9.26   1.99   4.34   196650   10.91   2.52   8.15   4.78   2.95   2.51   8.35   3.04   4.32   1814720   1.46   0.34   3.52   5.00   4.19   4.54   4.37   4.32   1841220   1.46   0.34   3.52   5.00   4.19   4.54   4.37   4.32   184733   15.01   3.48   3.75   5.14   4.04   6.05   2.61   4.32   184733   15.01   3.48   3.75   5.14   4.04   6.05   2.61   4.32   185650   14.94   3.47   5.20   2.99   5.04   3.92   4.39   4.31   155314   1.52   0.35   7.93   5.83   2.29   2.55   2.92   4.30   1417908   12.10   2.82   8.40   4.93   3.98   1.81   2.34   4.29   29063   3.34   0.78   2.94   6.06   4.11   3.85   4.48   4.29   2773345   1.21   0.28   3.42   4.56   2.89   8.45   2.12   4.29   18592006   5.30   1.24   4.98   3.66   4.06   4.48   4.22   4.28   461499   2.26   0.53   2.94   3.53   6.30   7.11   1.48   4.27   427838   4.05   0.95   7.99   6.36   2.29   2.32   2.29   4.25   1815800   1.22   0.29   5.36   6.04   5.43   3.31   1.13   4.25   1815801   1.22   0.29   5.36   6.04   5.43   3.31   1.13   4.25   1816409   1.22   0.29   5.36   6.04   5.43   3.31   1.13   4.25   1818404   4.04   0.95   3.84   7.52   3.07   4.21   2.54   1818444   4.04   0.95   3.84   7.52   3.07   4.21   2.54   1818444   4.04   0.95   3.84   7.52   3.07   4.21   2.54   1818449   2.501   5.91   4.16   5.55   3.42   4.66   3.40   4.22   1838850   2.29   3.39   3.81   2.13   4.52   1838800   6.77   1.62   4.54   3.37   5.89   5.18   1.97   4.19   182737   1.10   4.07   3.83   3.50   3.92   3.90   5.86   4.20   1828482   2.20   0.53   3.81   7.52   3.07   4.21   2.54   4.24   1831849   2.501   5.91   4.16   5.55   3.42   4.66   2.60   3.40   4.22   1838850   3.40   0.95   3.84   7.52   3.07   4.21   2.54   4.24   1831849   3.60   3.89   3.79   5.31   3.37   4.90   4.22   1831850   4.89   3.60   6.06   6.06   6.06   6.07   6.02   6.06   6.07   6.02   6.06   6.07   6.02   6.		•	1.46	1.85	6.04	4.99	6.43	2.63	
121436   6.37   1.46   7.91   6.01   2.52   3.24   2.09   4.36   814798   32.86   7.58   3.15   4.62   2.68   9.26   1.99   4.34   196650   10.93   2.52   4.17   6.60   2.20   6.71   1.99   4.34   196650   10.93   2.52   4.17   6.60   2.20   6.71   1.99   4.34   4.37   4.32   200599   13.42   3.11   5.88   2.44   8.89   1.51   2.87   4.32   200599   13.42   3.11   5.88   2.44   8.89   1.51   2.87   4.32   200599   13.42   3.11   5.88   2.44   8.89   1.51   2.87   4.32   200599   13.42   3.11   5.88   2.44   8.89   1.51   2.87   4.32   200599   13.42   3.11   5.88   2.44   8.89   1.51   2.87   4.32   200599   13.42   3.11   5.88   2.44   8.89   1.51   2.87   4.32   200599   13.42   3.17   5.88   5.29   4.04   4.38   4.31   3.55   5.50   4.94   4.94   4.38   4.31   4.52   5.3314   1.52   0.35   7.93   5.83   2.29   2.55   2.92   4.30   4.39   4.31   4.52   4.30   4.39   4.31   4.52   4.30   4.39   4.31   4.52   4.30   4.39   4.31   4.52   4.30   4.39   4.31   4.52   4.30   4.39   4.31   4.52   4.30   4.39   4.31   4.52   4.30   4.39   4.31   4.52   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4.30   4			6.05	2.45	3.37	5.47	1.22		
238821         6.54         1.50         5.59         1.57         4.76         0.92         8.90         4.35           814798         32.86         7.58         3.15         4.62         2.268         9.26         1.99         4.33           971367         35.22         8.15         4.78         2.95         2.51         8.35         3.04         4.32           200599         13.42         3.11         5.88         2.44         8.89         1.51         2.87         4.32           487733         15.01         3.48         3.75         5.14         4.04         6.05         2.61         4.32           5664537         1.79         0.42         2.77         5.08         5.29         4.04         4.38         4.31           50606         14.94         3.47         5.20         2.99         5.04         3.92         4.39         4.31           253314         1.52         0.35         7.93         5.83         2.29         2.55         2.92         4.30           417908         12.10         2.82         8.40         4.93         3.98         1.81         2.34         4.29           29063         3.34	121436	6.37	1.46	7.91	6.01		3.24		
814798         32.86         7.58         3.15         4.62         2.68         9.26         1.99         4.34           196650         10.93         2.52         4.17         6.60         2.20         6.71         1.99         4.33           971367         35.22         8.15         4.78         2.95         2.51         8.35         3.04         4.32           841220         1.46         0.34         3.52         5.00         4.19         4.54         4.37         4.32           200599         13.42         3.11         5.88         2.44         8.89         1.51         2.87         4.32           564537         1.79         0.42         2.77         5.08         5.29         4.04         4.38         4.31           253314         1.52         0.35         7.93         5.83         2.29         2.55         2.92         4.30           417908         12.10         2.82         8.40         4.93         3.98         1.81         2.34         4.29           1592006         5.30         1.24         4.98         3.66         4.06         4.11         3.85         4.84         4.29           1592006 <td< td=""><td>238821</td><td>6.54</td><td>1.50</td><td>5.59</td><td>1.57</td><td></td><td></td><td>1</td><td></td></td<>	238821	6.54	1.50	5.59	1.57			1	
19650   10.93	814798	32.86	7.58	3.15	4.62				
971367  35.22  8.15  4.78  2.95  2.51  8.35  3.04  4.32  841220  1.46  0.34  3.52  5.00  4.19  4.54  4.37  4.32  487733  15.01  3.48  3.75  5.14  4.04  6.05  2.61  4.32  487733  15.01  3.48  3.75  5.14  4.04  6.05  2.61  4.32  564537  1.79  0.42  2.77  5.08  5.29  4.04  4.38  4.31  50606  14.94  3.47  5.20  2.99  5.04  3.92  4.39  4.31  253314  1.52  0.35  7.93  5.83  2.29  2.55  2.92  4.30  417908  12.10  2.82  8.40  4.93  3.98  1.81  2.34  4.29  29063  3.34  0.78  2.94  6.06  4.11  3.85  4.48  4.29  2773345  1.21  0.28  3.42  4.56  2.89  8.45  2.12  4.29  1592006  5.30  1.24  4.98  3.66  4.06  4.48  4.22  4.28  461499  2.26  0.53  2.94  3.53  6.30  7.11  1.48  4.27  815800  1.22  0.29  5.36  6.04  5.43  3.31  1.13  4.25  436769  168.72  39.68  3.44  3.03  2.92  5.33  6.54  4.25  427838  4.05  0.95  7.99  6.36  2.29  2.32  2.29  4.25  50188  10.75  2.53  4.92  4.45  4.01  3.41  4.46  4.25  1519515  0.85  10.75  2.53  4.92  4.45  4.01  3.41  4.46  4.25  1519515  0.85  0.20  3.89  3.77  5.31  3.37  4.90  4.25  138444  4.04  0.95  3.84  7.52  3.07  4.21  2.54  4.24  1631849  25.01  5.91  4.16  5.55  3.42  4.66  3.40  4.24  1631849  25.01  5.91  4.16  5.55  3.42  4.66  3.40  4.24  1631839  25.01  5.91  4.16  5.55  3.42  4.66  3.40  4.24  1631839  7.10  0.25  3.02  4.22  3.83  5.08  4.94  4.22  290378  17.10  4.07  3.83  3.50  3.92  3.90  5.86  4.20  290378  17.10  4.07  3.83  3.50  3.92  3.90  5.86  4.20  290378  17.10  4.07  3.83  3.50  3.92  3.90  5.86  4.20  290379  17.74  4.26  6.37  2.52  9.06  1.63  1.24  4.17  593929  17.74  4.26  6.37  2.52  9.06  1.63  1.24  4.17  587415  12.98  3.12  7.46  3.53  7.08  1.19  1.53  4.16  886538  7.64  1.86  3.08  5.14  3.35  5.78  2.66  4.18  886914  18.03  4.35  5.29  7.59  0.53  6.00  1.32  4.11  1538414  4.20  1.99  0.59  1.24  9.68  5.69  3.37  4.11  1538414  8.20  1.99  0.59  1.24  9.68  5.69  3.37  4.11  1538414  8.20  1.99  0.59  1.24  9.68  5.69  3.37  4.11  1538414  1.52  2.81  1.99  0.59  1.24  9.68  5.69  3.37  4.11  1539399  4.22  1.03  4.35  5.29  4.75  9.06	196650	10.93	2.52	4.17					
841220         1.46         0.34         3.52         5.00         4.19         4.54         4.37         4.32           200599         13.42         3.11         5.88         2.44         8.89         1.51         2.87         4.32           487733         15.01         3.48         3.75         5.14         4.04         6.05         2.61         4.32           564537         1.79         0.42         2.77         5.08         5.29         4.04         4.38         4.31           50606         14.94         3.47         5.20         2.99         5.04         3.92         4.39         4.31           253314         1.52         0.35         7.93         5.83         2.29         2.55         2.92         4.30           417908         12.10         2.82         8.40         4.93         3.98         1.81         2.34         4.29           29063         3.34         0.78         2.94         6.06         4.11         3.85         4.48         4.29           1592006         5.30         1.24         4.98         3.66         4.06         4.48         4.22         4.28           451890         1.22         0.2	971367	35.22	8.15	4.78					
2005999         13.42         3.11         5.88         2.44         8.89         1.51         2.87         4.32           487733         15.01         3.48         3.75         5.14         4.04         6.05         2.61         4.32           5664537         1.79         0.42         2.77         5.08         5.29         4.04         4.38         4.31           253314         1.52         0.35         7.93         5.83         2.29         2.55         2.92         4.39         4.31           253314         1.52         0.35         7.93         5.83         2.29         2.55         2.92         4.30         4.21         4.29           29063         3.34         0.78         2.94         6.06         4.11         3.85         4.48         4.29           29063         3.34         0.78         2.94         4.56         2.89         8.45         2.12         4.29         4.26           461499         2.26         0.53         2.94         3.53         6.30         7.11         1.48         4.27           815800         1.22         3.95.68         3.44         3.03         2.92         5.33         6.54	841220	1.46	0.34						
487733         15.01         3.48         3.75         5.14         4.04         6.05         2.61         4.32           564537         1.79         0.42         2.77         5.08         5.29         4.04         4.38         4.31           50606         14.94         3.47         5.20         2.99         5.04         3.92         4.39         4.31           253314         1.52         0.35         7.93         5.83         2.29         2.55         2.92         4.30           417908         12.10         2.82         8.40         4.93         3.98         1.81         2.34         4.29           29063         3.34         0.78         2.94         6.06         4.11         3.85         4.48         4.29           773345         1.21         0.28         3.42         4.56         2.89         8.45         2.12         4.29           461499         2.26         0.53         2.94         3.53         6.30         7.11         1.48         4.27           415800         1.22         4.26         0.53         2.94         3.53         6.30         7.11         1.48         4.27           427838         4.05<	200599	13.42	3.11					1	
564537   1.79         0.42         2.77         5.08         5.29         4.04         4.38         4.31           50606   14.94   3.47   5.20         2.99         5.04   3.92   4.39         4.31         4.31           253314   1.52   0.35   7.93   5.83   2.29   2.55   2.92   4.30         417908   12.10   2.82   8.40   4.93   3.98   1.81   2.34   4.29         29063   3.34   0.78   2.94   6.06   4.11   3.85   4.48   4.29         4.29           29063   3.34   0.78   2.94   6.06   4.11   3.85   4.48   4.29   4.29         461499   6.66   4.06   4.48   4.22   4.29         4.29           1592006   5.30   1.24   4.98   3.66   6.06   4.06   4.48   4.22   4.28   4.04   4.98   4.06   4.48   4.22   4.28   4.04   4.95   4.05   4.25   4.04   4.25   4.04   4.25   4.04   4.25   4.04   4.25   4.04   4.25   4.04   4.25   4.04   4.25   4.04   4.25   4.04   4.25   4.04   4.25   4.04   4.05   4.25   4.25   4.24   4.04   4.05   0.95   7.99   6.36   2.29   2.32   2.29   4.25   4.25   4.24   4.04   4.05   4.25   4.24   4.04   4.05   4.25   4.24   4.04   4.05   4.25   4.24   4.04   4.05   4.25   4.24   4.04   4.05   4.25   4.24   4.04   4.05   4.24   4.04   4.05   4.24   4.04   4.05   4.24   4.04   4.05   4.24   4.04   4.05   4.25   4.24   4.04   4.05   4.24   4.04   4.05   4.25   4.24   4.04   4.05   4.25   4.24   4.04   4.07   4.07   3.83   3.50   3.92   3.90   5.86   4.20   4.24   4.04   4.07   3.83   3.50   3.92   3.90   5.86   4.20   4.24   4.04   4.07   4.04   4.22   4.25   4.25   4.24   4.04   4.25   4.24   4.05   4.25   4.24   4.05   4.25   4.24   4.05   4.25   4.24   4.05   4.25   4.25   4.25   4.25   4.25   4.25   4.25   4.25   4.25   4.25   4.25   4.25   4.25   4.25   4.25   4.25   4.25   4.25   4.25   4.25   4.25   4.25   4.25   4.25   4.25   4.25   4.25	487733	15.01	3.48						
50606         14.94         3.47         5.20         2.99         5.04         3.92         4.39         4.31           253314         1.52         0.35         7.93         5.83         2.29         2.55         2.92         4.30           417908         12.10         2.82         8.40         4.93         3.98         1.81         2.34         4.29           29063         3.34         0.78         2.94         6.06         4.11         3.85         4.48         4.29           173345         1.21         0.28         3.42         4.56         2.89         8.45         2.12         4.29           1592006         5.30         1.24         4.98         3.66         4.06         4.48         4.22         4.28           461499         2.26         0.53         2.94         3.53         6.30         7.11         1.48         4.27           815800         1.22         0.29         5.36         6.04         5.43         3.31         1.13         4.25           427838         4.05         0.95         7.99         6.36         2.29         2.32         2.29         4.25           5191515         0.82         0.20	564537	1.79						•	
253314         1.52         0.35         7.93         5.83         2.29         2.55         2.92         4.30           417908         12.10         2.82         8.40         4.93         3.98         1.81         2.34         4.29           29063         3.34         0.78         2.94         6.06         4.11         3.85         4.48         4.29           1592006         5.30         1.24         4.98         3.66         4.06         4.48         4.22         4.28           461499         2.26         0.53         2.94         3.53         6.30         7.11         1.48         4.27           436769         168.72         39.68         3.44         3.03         2.92         5.33         6.54         4.25           427838         4.05         0.95         7.99         6.36         2.29         2.32         2.29         4.25           1519515         0.85         0.20         3.89         3.77         5.31         3.37         4.90         4.25           1519515         0.85         0.20         3.89         3.77         5.31         3.37         4.90         4.25           1519517         1.07 <td< td=""><td>50606</td><td>14.94</td><td></td><td></td><td></td><td></td><td></td><td></td><td></td></td<>	50606	14.94							
417908         12.10         2.82         8.40         4.93         3.98         1.81         2.34         4.29           29063         3.34         0.78         2.94         6.06         4.11         3.85         4.48         4.29           173345         1.21         0.28         3.42         4.56         2.89         8.45         2.12         4.29           1592006         5.30         1.24         4.98         3.66         4.06         4.48         4.22         4.28           461499         2.26         0.53         2.94         3.53         6.30         7.11         1.48         4.27           815800         1.22         0.29         5.36         6.04         5.43         3.31         1.13         4.25           427838         4.05         0.95         7.99         6.36         2.29         2.32         2.29         4.25           50188         10.75         2.53         4.92         4.45         4.01         3.41         4.46         4.25           1519515         0.85         0.20         3.89         3.77         5.31         3.37         4.90         4.25           1519444         4.04         0.9	253314	1.52							
29063         3.34         0.78         2.94         6.06         4.11         3.85         4.48         4.29           773345         1.21         0.28         3.42         4.56         2.89         8.45         2.12         4.29           1592006         5.30         1.24         4.98         3.66         4.06         4.48         4.22         4.28           461499         2.26         0.53         3.53         6.30         7.11         1.48         4.27           815800         1.22         0.29         5.36         6.04         5.43         3.31         1.13         4.25           427838         4.05         0.95         7.99         6.36         2.29         2.32         2.29         4.25           427838         4.05         0.95         7.99         6.36         2.29         2.32         2.29         4.25           1519515         0.85         0.20         3.89         3.77         5.31         3.37         4.90         4.25           1519515         0.85         0.20         3.84         7.52         3.07         4.21         2.54         4.24           1631849         25.01         4.16         5.									
773345         1.21         0.28         3.42         4.56         2.89         8.45         2.12         4.29           1592006         5.30         1.24         4.98         3.66         4.06         4.48         4.22         4.28           461499         2.26         0.53         2.94         3.53         6.30         7.11         1.48         4.27           815800         1.22         0.29         5.36         6.04         5.43         3.31         1.13         4.25           436769         168.72         39.68         3.44         3.03         2.92         5.33         6.54         4.25           427838         4.05         0.95         7.99         6.36         2.29         2.32         2.29         4.25           50188         10.75         2.53         4.92         4.45         4.01         3.41         4.46         4.25           1519515         0.85         0.20         3.89         3.77         5.31         3.31         4.90         4.25           1631849         25.01         5.91         4.16         5.55         3.42         4.66         3.40         4.24           613237         1.07 <td< td=""><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td></td<>									
1592006         5.30         1.24         4.98         3.66         4.06         4.48         4.22         428           461499         2.26         0.53         2.94         3.53         6.30         7.11         1.48         4.27           815800         1.22         0.29         5.36         6.04         5.43         3.31         1.13         4.25           436769         168.72         39.68         3.44         3.03         2.92         5.33         6.54         4.25           427838         4.05         0.95         7.99         6.36         2.29         2.32         2.29         4.25           50188         10.75         2.53         4.92         4.45         4.01         3.41         4.46         4.25           1519515         0.85         0.20         3.89         3.77         5.31         3.37         4.90         4.25           153844         4.04         0.95         3.84         7.52         3.07         4.21         2.54         4.24           1631849         25.01         5.91         4.16         5.55         3.42         4.66         3.40         4.24           613227         1.07									
461499         2.26         0.53         2.94         3.53         6.30         7.11         1.48         4.27           815800         1.22         0.29         5.36         6.04         5.43         3.31         1.13         4.25           436769         168.72         39.68         3.44         3.03         2.92         5.33         6.54         4.25           427838         4.05         0.95         7.99         6.36         2.29         2.32         2.29         4.25           50188         10.75         2.53         4.92         4.45         4.01         3.41         4.46         4.25           1519515         0.85         0.20         3.89         3.77         5.31         3.37         4.90         4.25           1631849         25.01         5.91         4.16         5.55         3.42         4.66         3.40         4.24           1631237         1.07         0.25         3.02         4.22         3.83         5.08         4.94         4.22           358850         21.29         5.06         7.24         3.98         3.18         2.13         4.52         4.21           290378         17.10         <									
815800         1.22         0.29         5.36         6.04         5.43         3.31         1.13         4.25           436769         168.72         39.68         3.44         3.03         2.92         5.33         6.54         4.25           427838         4.05         0.95         7.99         6.36         2.29         2.32         2.29         4.25           50188         10.75         2.53         4.92         4.45         4.01         3.41         4.46         4.25           1519515         0.85         0.20         3.89         3.77         5.31         3.37         4.99         4.25           138444         4.04         0.95         3.84         7.52         3.07         4.21         2.54         4.24           1631849         25.01         5.91         4.16         5.55         3.42         4.66         3.40         4.24           613237         1.07         0.25         3.02         4.22         3.83         5.08         4.94         4.22           290378         17.10         4.07         3.83         3.50         3.92         3.90         5.86         4.20           898109         6.77 <td< td=""><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td></td<>									
436769         168.72         39.68         3.44         3.03         2.92         5.33         6.54         4.25           427838         4.05         0.95         7.99         6.36         2.29         2.32         2.29         4.25           50188         10.75         2.53         4.92         4.45         4.01         3.41         4.46         4.25           1519515         0.85         0.20         3.89         3.77         5.31         3.37         4.90         4.25           138444         4.04         0.95         3.84         7.52         3.07         4.21         2.54         4.24           6131849         25.01         5.91         4.16         5.55         3.42         4.66         3.40         4.24           613237         1.07         0.25         3.02         4.22         3.83         5.08         4.94         4.22           358850         21.29         5.06         7.24         3.98         3.18         2.13         4.52         4.21           898109         6.77         1.62         4.54         3.37         5.89         5.18         1.97         4.19           82734         7.21									
427838         4.05         0.95         7.99         6.36         2.29         2.32         2.29         4.25           50188         10.75         2.53         4.92         4.45         4.01         3.41         4.46         4.25           1519515         0.85         0.20         3.89         3.77         5.31         3.37         4.90         4.25           138444         4.04         0.95         3.84         7.52         3.07         4.21         2.54         4.24           1631849         25.01         5.91         4.16         5.55         3.42         4.66         3.40         4.24           613237         1.07         0.25         3.02         4.22         3.83         5.08         4.94         4.22           358850         21.29         5.06         7.24         3.98         3.18         2.13         4.52         4.21           290378         17.10         4.07         3.83         3.50         3.92         3.90         5.86         4.20           88109         6.77         1.62         4.54         3.37         5.89         5.18         1.97         4.19           82734         7.21         1.7									
50188         10.75         2.53         4.92         4.45         4.01         3.41         4.46         4.25           1519515         0.85         0.20         3.89         3.77         5.31         3.37         4.90         4.25           138444         4.04         0.95         3.84         7.52         3.07         4.21         2.54         4.24           1631849         25.01         5.91         4.16         5.55         3.42         4.66         3.40         4.24           613237         1.07         0.25         3.02         4.22         3.83         5.08         4.94         4.22           358850         21.29         5.06         7.24         3.98         3.18         2.13         4.52         4.21           290378         17.10         4.07         3.83         3.50         3.92         3.90         5.86         4.20           898109         6.77         1.62         4.54         3.37         5.89         5.18         1.97         4.19           82734         7.21         1.73         3.01         4.42         4.26         6.62         2.60         4.18           450307         9.39         2.									
1519515         0.85         0.20         3.89         3.77         5.31         3.37         4.90         4.25           138444         4.04         0.95         3.84         7.52         3.07         4.21         2.54         4.24           1631849         25.01         5.91         4.16         5.55         3.42         4.66         3.40         4.24           613237         1.07         0.25         3.02         4.22         3.83         5.08         4.94         4.22           358850         21.29         5.06         7.24         3.98         3.18         2.13         4.52         4.21           290378         17.10         4.07         3.83         3.50         3.92         3.90         5.86         4.20           898109         6.77         1.62         4.54         3.37         5.89         5.18         1.97         4.19           82734         7.21         1.73         3.01         4.42         4.26         6.62         2.60         4.18           450307         9.39         2.25         7.70         6.02         3.75         1.27         2.10         4.17           587415         12.98         3									
138444         4.04         0.95         3.84         7.52         3.07         4.21         2.54         4.24           1631849         25.01         5.91         4.16         5.55         3.42         4.66         3.40         4.24           613237         1.07         0.25         3.02         4.22         3.83         5.08         4.94         4.22           358850         21.29         5.06         7.24         3.98         3.18         2.13         4.52         4.21           290378         17.10         4.07         3.83         3.50         3.92         3.90         5.86         4.20           898109         6.77         1.62         4.54         3.37         5.89         5.18         1.97         4.19           82734         7.21         1.73         3.01         4.42         4.26         6.62         2.60         4.18           450307         9.39         2.25         7.70         6.02         3.75         1.27         2.10         4.17           587415         12.98         3.12         7.46         3.53         7.08         1.19         1.53         4.16           842882         2.20         0.									
1631849         25.01         5.91         4.16         5.55         3.42         4.66         3.40         4.24           613237         1.07         0.25         3.02         4.22         3.83         5.08         4.94         4.22           358850         21.29         5.06         7.24         3.98         3.18         2.13         4.52         4.21           290378         17.10         4.07         3.83         3.50         3.92         3.90         5.86         4.20           898109         6.77         1.62         4.54         3.37         5.89         5.18         1.97         4.19           82734         7.21         1.73         3.01         4.42         4.26         6.62         2.60         4.18           450307         9.39         2.25         7.70         6.02         3.75         1.27         2.10         4.17           587415         12.98         3.12         7.46         3.53         7.08         1.19         1.53         4.16           842882         2.20         0.53         3.81         5.06         5.56         2.68         3.67         4.16           896914         18.03         4	138444	4.04							
613237         1.07         0.25         3.02         4.22         3.83         5.08         4.94         4.22           358850         21.29         5.06         7.24         3.98         3.18         2.13         4.52         4.21           290378         17.10         4.07         3.83         3.50         3.92         3.90         5.86         4.20           898109         6.77         1.62         4.54         3.37         5.89         5.18         1.97         4.19           82734         7.21         1.73         3.01         4.42         4.26         6.62         2.60         4.18           450307         9.39         2.25         7.70         6.02         3.75         1.27         2.10         4.17           587415         12.98         3.12         7.46         3.53         7.08         1.19         1.53         4.16           842882         2.20         0.53         3.81         5.06         5.56         2.68         3.67         4.16           896914         18.03         4.35         5.29         7.59         0.53         6.00         1.32         4.14           1607765         3.96         0.									
358850         21.29         5.06         7.24         3.98         3.18         2.13         4.52         4.21           290378         17.10         4.07         3.83         3.50         3.92         3.90         5.86         4.20           898109         6.77         1.62         4.54         3.37         5.89         5.18         1.97         4.19           82734         7.21         1.73         3.01         4.42         4.26         6.62         2.60         4.18           450307         9.39         2.25         7.70         6.02         3.75         1.27         2.10         4.17           587415         12.98         3.12         7.46         3.53         7.08         1.19         1.53         4.16           842882         2.20         0.53         3.81         5.06         5.56         2.68         3.67         4.16           896914         18.03         4.35         5.29         7.59         0.53         6.00         1.32         4.14           1607765         3.96         0.96         4.88         3.54         3.13         5.17         3.98         4.14           824329         1.94         0.		1							
290378         17.10         4.07         3.83         3.50         3.92         3.90         5.86         4.20           898109         6.77         1.62         4.54         3.37         5.89         5.18         1.97         4.19           82734         7.21         1.73         3.01         4.42         4.26         6.62         2.60         4.18           450307         9.39         2.25         7.70         6.02         3.75         1.27         2.10         4.17           593929         17.74         4.26         6.37         2.52         9.06         1.63         1.24         4.17           587415         12.98         3.12         7.46         3.53         7.08         1.19         1.53         4.16           842882         2.20         0.53         3.81         5.06         5.56         2.68         3.67         4.16           89614         18.03         4.35         5.29         7.59         0.53         6.00         1.32         4.14           1607765         3.96         0.96         4.88         3.54         3.13         5.17         3.98         4.14           824329         1.94         0.4	358850	21.29	5.06						
898109         6.77         1.62         4.54         3.37         5.89         5.18         1.97         4.19           82734         7.21         1.73         3.01         4.42         4.26         6.62         2.60         4.18           450307         9.39         2.25         7.70         6.02         3.75         1.27         2.10         4.17           593929         17.74         4.26         6.37         2.52         9.06         1.63         1.24         4.17           587415         12.98         3.12         7.46         3.53         7.08         1.19         1.53         4.16           842882         2.20         0.53         3.81         5.06         5.56         2.68         3.67         4.16           896914         18.03         4.35         5.29         7.59         0.53         6.00         1.32         4.14           1607765         3.96         0.96         4.88         3.54         3.13         5.17         3.98         4.14           824329         1.94         0.47         4.04         4.22         5.97         3.83         2.60         4.13           1558411         8.20         1.	290378	17.10	4.07	3.83					
82734         7.21         1.73         3.01         4.42         4.26         6.62         2.60         4.18           450307         9.39         2.25         7.70         6.02         3.75         1.27         2.10         4.17           593929         17.74         4.26         6.37         2.52         9.06         1.63         1.24         4.17           587415         12.98         3.12         7.46         3.53         7.08         1.19         1.53         4.16           842882         2.20         0.53         3.81         5.06         5.56         2.68         3.67         4.16           896914         18.03         4.35         5.29         7.59         0.53         6.00         1.32         4.14           1607765         3.96         0.96         4.88         3.54         3.13         5.17         3.98         4.14           824329         1.94         0.47         4.04         4.22         5.97         3.83         2.60         4.13           1558411         8.20         1.99         0.59         1.24         9.68         5.69         3.37         4.11           485359         4.23         1.	898109	6.77	1.62	4.54					
450307         9.39         2.25         7.70         6.02         3.75         1.27         2.10         4.17           593929         17.74         4.26         6.37         2.52         9.06         1.63         1.24         4.17           587415         12.98         3.12         7.46         3.53         7.08         1.19         1.53         4.16           842882         2.20         0.53         3.81         5.06         5.56         2.68         3.67         4.16           896914         18.03         4.35         5.29         7.59         0.53         6.00         1.32         4.14           1607765         3.96         0.96         4.88         3.54         3.13         5.17         3.98         4.14           824329         1.94         0.47         4.04         4.22         5.97         3.83         2.60         4.13           1558411         8.20         1.99         0.59         1.24         9.68         5.69         3.37         4.11           453599         4.23         1.03         3.89         3.56         3.50         6.56         3.01         4.11           486538         7.64         1	82734	7.21	1.73	3.01					
593929         17.74         4.26         6.37         2.52         9.06         1.63         1.24         4.17           587415         12.98         3.12         7.46         3.53         7.08         1.19         1.53         4.16           842882         2.20         0.53         3.81         5.06         5.56         2.68         3.67         4.16           896914         18.03         4.35         5.29         7.59         0.53         6.00         1.32         4.14           1607765         3.96         0.96         4.88         3.54         3.13         5.17         3.98         4.14           824329         1.94         0.47         4.04         4.22         5.97         3.83         2.60         4.13           1558411         8.20         1.99         0.59         1.24         9.68         5.69         3.37         4.11           453599         4.23         1.03         3.89         3.56         3.50         6.56         3.01         4.11           486538         7.64         1.86         3.08         5.14         3.85         5.78         2.65         4.10           712216         0.50         0	450307	9.39	2.25	7.70	6.02				
587415         12.98         3.12         7.46         3.53         7.08         1.19         1.53         4.16           842882         2.20         0.53         3.81         5.06         5.56         2.68         3.67         4.16           896914         18.03         4.35         5.29         7.59         0.53         6.00         1.32         4.14           1607765         3.96         0.96         4.88         3.54         3.13         5.17         3.98         4.14           824329         1.94         0.47         4.04         4.22         5.97         3.83         2.60         4.13           1558411         8.20         1.99         0.59         1.24         9.68         5.69         3.37         4.11           453599         4.23         1.03         3.89         3.56         3.50         6.56         3.01         4.11           486538         7.64         1.86         3.08         5.14         3.85         5.78         2.65         4.10           712216         0.50         0.12         5.35         4.81         4.78         4.53         1.03         4.10           753162         2.81         0.	593929	17.74	4.26	6.37	2.52				
842882         2.20         0.53         3.81         5.06         5.56         2.68         3.67         4.16           896914         18.03         4.35         5.29         7.59         0.53         6.00         1.32         4.14           1607765         3.96         0.96         4.88         3.54         3.13         5.17         3.98         4.14           824329         1.94         0.47         4.04         4.22         5.97         3.83         2.60         4.13           1558411         8.20         1.99         0.59         1.24         9.68         5.69         3.37         4.11           453599         4.23         1.03         3.89         3.56         3.50         6.56         3.01         4.11           486538         7.64         1.86         3.08         5.14         3.85         5.78         2.65         4.10           712216         0.50         0.12         5.35         4.81         4.78         4.53         1.03         4.10           1637296         25.52         6.23         4.17         2.86         2.82         6.06         4.57         4.09           898242         10.34	587415	12.98	3.12	7.46	3.53				
896914         18.03         4.35         5.29         7.59         0.53         6.00         1.32         4.14           1607765         3.96         0.96         4.88         3.54         3.13         5.17         3.98         4.14           824329         1.94         0.47         4.04         4.22         5.97         3.83         2.60         4.13           1558411         8.20         1.99         0.59         1.24         9.68         5.69         3.37         4.11           453599         4.23         1.03         3.89         3.56         3.50         6.56         3.01         4.11           486538         7.64         1.86         3.08         5.14         3.85         5.78         2.65         4.10           712216         0.50         0.12         5.35         4.81         4.78         4.53         1.03         4.10           753162         2.81         0.69         2.94         3.35         2.20         9.53         2.46         4.10           1637296         25.52         6.23         4.17         2.86         2.82         6.06         4.57         4.09           898242         10.34	842882	2.20	0.53	3.81	5.06	5.56		3.67	
1607765         3.96         0.96         4.88         3.54         3.13         5.17         3.98         4.14           824329         1.94         0.47         4.04         4.22         5.97         3.83         2.60         4.13           1558411         8.20         1.99         0.59         1.24         9.68         5.69         3.37         4.11           453599         4.23         1.03         3.89         3.56         3.50         6.56         3.01         4.11           486538         7.64         1.86         3.08         5.14         3.85         5.78         2.65         4.10           712216         0.50         0.12         5.35         4.81         4.78         4.53         1.03         4.10           753162         2.81         0.69         2.94         3.35         2.20         9.53         2.46         4.10           1637296         25.52         6.23         4.17         2.86         2.82         6.06         4.57         4.09           898242         10.34         2.53         2.92         4.71         4.80         5.15         2.87         4.09           771048         10.54	896914	18.03	4.35	5.29	7.59	0.53			
824329         1.94         0.47         4.04         4.22         5.97         3.83         2.60         4.13           1558411         8.20         1.99         0.59         1.24         9.68         5.69         3.37         4.11           453599         4.23         1.03         3.89         3.56         3.50         6.56         3.01         4.11           486538         7.64         1.86         3.08         5.14         3.85         5.78         2.65         4.10           712216         0.50         0.12         5.35         4.81         4.78         4.53         1.03         4.10           753162         2.81         0.69         2.94         3.35         2.20         9.53         2.46         4.10           1637296         25.52         6.23         4.17         2.86         2.82         6.06         4.57         4.09           898242         10.34         2.53         2.92         4.71         4.80         5.15         2.87         4.09           771048         10.54         2.58         5.50         3.59         3.73         3.20         4.39         4.08           951125         18.86	1607765	3.96	0.96	4.88	3.54	3.13		3.98	
1558411         8.20         1.99         0.59         1.24         9.68         5.69         3.37         4.11           453599         4.23         1.03         3.89         3.56         3.50         6.56         3.01         4.11           486538         7.64         1.86         3.08         5.14         3.85         5.78         2.65         4.10           712216         0.50         0.12         5.35         4.81         4.78         4.53         1.03         4.10           753162         2.81         0.69         2.94         3.35         2.20         9.53         2.46         4.10           1637296         25.52         6.23         4.17         2.86         2.82         6.06         4.57         4.09           898242         10.34         2.53         2.92         4.71         4.80         5.15         2.87         4.09           771048         10.54         2.58         5.50         3.59         3.73         3.20         4.39         4.08           951125         18.86         4.63         3.30         7.80         2.31         3.51         3.45         4.07           842925         4.61	824329	1.94	0.47	4.04	4.22	5.97	3.83	2.60	
453599         4.23         1.03         3.89         3.56         3.50         6.56         3.01         4.11           486538         7.64         1.86         3.08         5.14         3.85         5.78         2.65         4.10           712216         0.50         0.12         5.35         4.81         4.78         4.53         1.03         4.10           753162         2.81         0.69         2.94         3.35         2.20         9.53         2.46         4.10           1637296         25.52         6.23         4.17         2.86         2.82         6.06         4.57         4.09           898242         10.34         2.53         2.92         4.71         4.80         5.15         2.87         4.09           771048         10.54         2.58         5.50         3.59         3.73         3.20         4.39         4.08           951125         18.86         4.63         3.30         7.80         2.31         3.51         3.45         4.07           842925         4.61         1.13         3.78         3.80         4.94         4.96         2.87         4.07	1558411	8.20	1.99	0.59	1.24	9.68			
486538         7.64         1.86         3.08         5.14         3.85         5.78         2.65         4.10           712216         0.50         0.12         5.35         4.81         4.78         4.53         1.03         4.10           753162         2.81         0.69         2.94         3.35         2.20         9.53         2.46         4.10           1637296         25.52         6.23         4.17         2.86         2.82         6.06         4.57         4.09           898242         10.34         2.53         2.92         4.71         4.80         5.15         2.87         4.09           771048         10.54         2.58         5.50         3.59         3.73         3.20         4.39         4.08           951125         18.86         4.63         3.30         7.80         2.31         3.51         3.45         4.07           611150         7.99         1.96         4.60         2.92         4.15         5.04         3.66         4.07           842925         4.61         1.13         3.78         3.80         4.94         4.96         2.87         4.07	453599	4.23	1.03	3.89	3.56	3.50	6.56	3.01	
753162         2.81         0.69         2.94         3.35         2.20         9.53         2.46         4.10           1637296         25.52         6.23         4.17         2.86         2.82         6.06         4.57         4.09           898242         10.34         2.53         2.92         4.71         4.80         5.15         2.87         4.09           771048         10.54         2.58         5.50         3.59         3.73         3.20         4.39         4.08           951125         18.86         4.63         3.30         7.80         2.31         3.51         3.45         4.07           611150         7.99         1.96         4.60         2.92         4.15         5.04         3.66         4.07           842925         4.61         1.13         3.78         3.80         4.94         4.96         2.87         4.07			1.86	3.08	5.14	3.85		2.65	
1637296     25.52     6.23     4.17     2.86     2.82     6.06     4.57     4.09       898242     10.34     2.53     2.92     4.71     4.80     5.15     2.87     4.09       771048     10.54     2.58     5.50     3.59     3.73     3.20     4.39     4.08       951125     18.86     4.63     3.30     7.80     2.31     3.51     3.45     4.07       611150     7.99     1.96     4.60     2.92     4.15     5.04     3.66     4.07       842925     4.61     1.13     3.78     3.80     4.94     4.96     2.87     4.07	712216	0.50	0.12	5.35	4.81	4.78	4.53	1.03	4.10
898242     10.34     2.53     2.92     4.71     4.80     5.15     2.87     4.09       771048     10.54     2.58     5.50     3.59     3.73     3.20     4.39     4.08       951125     18.86     4.63     3.30     7.80     2.31     3.51     3.45     4.07       611150     7.99     1.96     4.60     2.92     4.15     5.04     3.66     4.07       842925     4.61     1.13     3.78     3.80     4.94     4.96     2.87     4.07				2.94	3.35	2.20	9.53	2.46	4.10
898242     10.34     2.53     2.92     4.71     4.80     5.15     2.87     4.09       771048     10.54     2.58     5.50     3.59     3.73     3.20     4.39     4.08       951125     18.86     4.63     3.30     7.80     2.31     3.51     3.45     4.07       611150     7.99     1.96     4.60     2.92     4.15     5.04     3.66     4.07       842925     4.61     1.13     3.78     3.80     4.94     4.96     2.87     4.07		1			2.86	2.82		4.57	4.09
771048     10.54     2.58     5.50     3.59     3.73     3.20     4.39     4.08       951125     18.86     4.63     3.30     7.80     2.31     3.51     3.45     4.07       611150     7.99     1.96     4.60     2.92     4.15     5.04     3.66     4.07       842925     4.61     1.13     3.78     3.80     4.94     4.96     2.87     4.07				2.92	4.71	4.80		2.87	4.09
951125     18.86     4.63     3.30     7.80     2.31     3.51     3.45     4.07       611150     7.99     1.96     4.60     2.92     4.15     5.04     3.66     4.07       842925     4.61     1.13     3.78     3.80     4.94     4.96     2.87     4.07				5.50	3.59				4.08
611150 7.99 1.96 4.60 2.92 4.15 5.04 3.66 4.07 842925 4.61 1.13 3.78 3.80 4.94 4.96 2.87 4.07				3.30	7.80				
842925 4.61 1.13 3.78 3.80 4.94 4.96 2.87 4.07				4.60	2.92				4.07
824843   4.03   0.99   5.28 3.89 3.11 6.61 1.41 4.06				3.78	3.80	4.94	4.96	2.87	4.07
	824843	4.03	0.99	5.28	3.89	3.11	6.61	1.41	4.06

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Table 1-1

878544	19.20	4.73	8.54	2.29	4.85	2.59	2.02	4.06
781366	12.06	2.98	4.50	3.42	1.61	2.86	7.87	4.05
79000	6.24	1.54	2.16	5.95	3.81	5.31	3.04	4.05
24208	7.49	1.85	3.19	3.67	3.57	4.54	5.28	4.05
729953	4.95	1.23	3.26	4.43	3.91	6.32	2.25	4.04
34134	2.73	0.68	6.48	3.68	4.71	3.19	2.08	4.03
745283	7.94	1.97	4.03	3.25	3.72	4.68	4.44	4.02
1602619	135.32	33.71	4.29	3.43	3.41	4.71	4.23	4.02
1636868	5.07	1.27	5.31	4.50	3.81	4.48	1.96	4.01
726483	3.50	0.87	2.71	4.78	2.55	5.48	4.52	4.01
360403	28.16	7.03	9.60	1.99	5.91	1.37	1.17	4.01
757462	12.77	3.19	4.05	4.75	4.11	4.73	2.41	4.01
279482	8.66	2.16	3.13	2.00	10.92	1.42	2.54	4.00
811766	4.93	1.23	1.80	3.53	4.42	6.47	3.79	4.00
50764	0.43	0.11	4.10	3.54	3.73	5.17	3.48	4.00
626716	21.60	5.41	2.78	6.84	3.77	3.86	2.73	4.00
213682	13.07	3.27	0.79	3.49	2.68	9.05	3.96	3.99
685922	3.59	0.90	3.85	3.96	3.20	5.02	3.93	3.99
1631194	21.83	5.47	5.50	1.73	5.19	6.07	1.46	3.99
823578	0.56	0.14	4.98	3.43	4.18	3.36	3.93	3.98
32509	4.38	1.10	1.50	7.60	1.95	4.07	4.72	3.97
845352	13.36	3.37	4.44	3.86	3.95	2.54	5.05	3.97
191978	47.95	12.11	3.93	4.27	3.76	5.36	2.48	3.96
120707	19.60	4.95	2.32	3 <b>.87</b>	3.65	8.32	1.62	3.96
75415	51.71	13.09	4.45	4.47	2.60	4.53	3.70	3.95
186768	9.43	2.39	2.81	1.73	11.34	2.18	1.69	3.95
80484	14.69	3.72	9.35	2.80	1.75	3.52	2.32	3.95
1475200	28.16	7.14	1.70	3.21	6.52	2.96	5.34	3.95
810510	27.71	7.03	3.76	3.64	2.68	4.64	5.00	3.94
897690	15.87	4.03	5.68	4.08	4.14	1.92	3.87	3.94
45327	8.18	2.08	2.60	3.75	4.04	5.11	4.18	3.94
299723	2.98	0.76	1.71	2.92	5.37	4.69	4.93	3.93
81578	12.77	3.25	5.02	3.62	6.26	2.36	2.37	3.93
197888	) 1	1.27	4.11	2.76	2.65	3.61	6.49	3.92
810571	4.37	1.11	2.29	5.21	2.51	6.03	3.57	3.92
768443		6.79	2.54	4.37	3 <b>.56</b>	7.05	2.07	3.92
150314		1.35	2.49	4.79	3.04	6.51	2.76	3.92
196189		2.86	4.68	4.67	3.98	2.71	3.52	3.91
626990		1.89	2.81	2.77	3.38	7.85	2.73	3.91
127396		3.61	3.73	5.03	1.41	6.75	2.59	3.90
49710		0.82	2.93	3.48	4.65	4.08	4.34	3.90
262864		9.52	3.65	2.59	3.11	5.73	4.36	3.89
815737		0.94	3.91	4.25	3.43	5.08	2.74	3.88
256680		0.44	3.22	3.59	3.08	4.91	4.60	3.88
461509		0.16	4.43	4.30	3.23	5.77	1.64	3.87
809578			5.73	2.71	3.10	4.48	3.32	3.87
81112			2.85	4.21	4.20	4.46	3.62	3.87
547058			4.03	2.83	3.49	4.83	4.11	3.86
103576			3.91	1.25	4.74	0.71	8.68	3.86
82473			7.31	4.22	2.34	3.52	1.89	3.86
81501	7 14.81	3.84	3.95	3.11	3.65	4.61	3.95	3.85

Table 1-1

32684	24.32	6.32	4.79	1.68	2.94	5.24	4.60	3.85
178137	238.41	61.93	3.66	2.98	2.56	6.38	3.66	3.85
282283	0.84	0.22	3 <b>.28</b>	3.36	3.61	4.89	4.09	3.85
725978	4.92	1.28	5.17	3.07	2.63	4.05	4.31	3.85
1631762	46.32	12.06	2.78	4.97	2.53	4.94	4.00	3.84
1553306	7.92	2.06	3.60	3.69	3.79	4.48	3.64	3.84
590145	4.37	1.14	3.78	3.14	3.77	5.80	2.69	3.84
815230	9.59	2.50	3.37	6.27	3.72	3.68	2.14	3.84
824237	74.89	19.54	3.14	3.18	3.55	5.72	3.57	3.83
361048	3.13	0.82	2.53	3.04	6.30	4.44	2.84	3.83
417867	13.52	3.54	2.81	3.70	3.15	4.96	4.47	3.82
878152	4.20	1.10	3.49	3.90	4.76	3.69	3.25	3.82
487499	4.41	1.16	2.68	3.09	8.83	3.10	1.37	3.82
712570	1.32	0.35	4.30	4.75	4.52	3.37	2.10	3.81
124824	30.85	8.13	3.79	2.43	2.17	3.37	7.23	3.80
626385	15.94	4.20	2.97	4.37	4.40	2.13	5.11	3.79
813271	0.96	0.25	3.16	4.24	3 <b>.70</b>	4.31	3.56	3.79
561971	3.81	1.01	2.50	4.39	4.74	3.83	3.46	3.79
824358	0.96	0.25	4.80	3.84	3.71	5.67	0.90	3.79
<b>7</b> 41067		17.00	0.88	4.25	2.30	7.0 <del>9</del>	4.38	3.78
46411		0.81	2.37	2.25	5.61	3.01	5.66	3.78
594517		6.57	4.24	1.83	7.82	1.45	3.53	3.78
825668		0.60	3.80	3.91	4.97	3.99	2.20	3.77
362926		1.17	1.58	4.23	3.24	2.08	7.71	3.77
814018		0.26	3.30	3.43	6.07	5.33	0.71	3.77
745503		11.32	2.92	3.95	3.33	3.03	5.59	3.77
884531		2.05	2.78	3.56	3.60	6.93	1.94	3.76
210862		0.68	5.33	3.36	3.68	3.66	2.76	3.76 3.76
504544		3.28	2.35	4.09	2.54	7.86	1.95 1.95	3.76 3.76
838668		0.95	1.88	5.28	2.47 2.56	7.20 6.57	3.78	3.74
433253		1.95 0.91	2.88 3.19	2.93 3.49	2.36 3.78	5.20	3.78	3.74
461330 87852:		2.97	4.03	4.61	3.7 <b>6</b> 3. <b>75</b>	2.79	3.49	3.74
38318		3.22	6.71	2.73	7.18	1.05	1.01	3.73
89773	1 1	1.58	4.19	4.06	4.13	4.55	1.60	3.71
159401		6.55	3.80	3.42	3.34	3.87	4.10	3.71
50656		3.49	5.12	3.30	3.70	2.28	4.10	3.70
82491	. 1		4.19	3.64	3.11	2.99	4.52	3.69
8495			5.70	4.55	4.49	1.79	1.92	3.69
95108			6.73	4.60	2.27	2.30	2.53	3.69
3042			0.76	1.72	2.50	10.04	3.40	3.68
78409			3.21	4.16	4.32	2.79	3.93	3.68
13590		1	9.57	1.87	4.30	1.31	1.34	3.68
95068			4.34	2.16	2.56	4.14	5.19	3.68
87828	4		5.36	2.63	2.49	1.62	6.22	3.66
84536			4.04	3.09	2.09	5.22	3.87	3.66
3584		1	4.06	4.52	2.15	4.94	2.64	3.66
762	1	1 1	5.18	3.55	3.64	3.20	2.69	3.65
366			4.19	3.70	2.34	5.99	2.02	3.65
6242	1		3.97	2.76	2.08	7.77	1.62	3.64
7430	71 1.89	9 0.52	4.55	1.78	1.17	8.79	1.87	3.63
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Table 1-1

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1070015	3.26	0.90	1.10	3.25	0.84	5.24	7.71	3.63
203132	12.63	3.48	2.95	2.71	2.51	4.70	5.27	3.63
259953	1.46	0.40	3.59	3.86	2.56	4.66	3.46	3.63
814427	0.47	0.13	3.78	3.80	3.96	5.65	0.91	3.62
259462	3.69	1.02	1.07	1.43	1.22	13.31	1.06	3.62
810558	14.49	4.01	2.38	3.34	3.00	3.69	5.66	3.61
51151	3.26	0.90	3.04	4.60	2.81	5.13	2.49	3.61
272750	20.57	5.70	3.98	5.64	2.94	2.81	2.70	3.61
360885	40.29	11.17	5.08	2.71	5.71	1.52	3.03	3.61
208082	7.90	2.19	2.97	6.19	3.18	2.98	2.71	3.61
627428	24.22	6.72	1.13	1.36	6.42	4.29	4.83	3.61
845345	5.09	1.41	2.54	3. <b>3</b> 2	3.32	4.11	4.73	3.61
417801	17.18	4.77	2.94	2.40	2.51	5.34	4.83	3.60
361922	9.18	2.55	3.94	3.22	3.20	3.45	4.16	3.60
898206	1.79	0.50	2.31	3.99	3.53	4.38	3.77	3.60
431296	1.85	0.51	2.75	3.55	5.68	1.98	4.01	3.59
587595	559.22	155.61	0.86	2.83	3.47	7.37	3.44	3.59
1632161	39.75	11.08	4.86	3.58	2.47	2.92	4.10	3.59
121196	22.80	6.36	2.64	3.37	3.11	3.53	5.26	3.58
823819	11.96	3.34	4.72	4.29	2.21	3.08	3.60	3.58
1456721	12.82	3.58	3.28	4.85	3.29	3.04	3.47	3.58
1500324	6.77	1.89	2.74	6.90	3.33	2.21	2.73	3.58
86160	2.10	0.59	3.16	4.11	3.48	3.18	3.96	3.58
246620	16.85	4.71	3.66	3.89	3.47	5.00	1.89	3.58
52730	6.16	1.72	2.03	3.11	2.60	6.50	3.64	3.58
809657	4.72	1.32	5.12	2.96	2.52	2.52	4.76	3.58
253 <b>577</b>	1.48	0.41	4.99	4.45	2.01	2.88	3.54	3.57
486459		1.75	2.96	3.80	3.21	5.49	2.40	3.57
23576	3.58	1.00	2.58	4.47	3.35	4.26	3.17	3.57
119403	1.22	0.34	3.95	4.15	2.67	3.43	3.62	3.57
324342		1.24	4.11	3.07	4.52	3.34	2.80	3.57
447518	10.98	3.08	1.93	4.33	4.07	2.68	4.81	3.56
811953		5.02	3 <b>.87</b>	4.29	4.39	2.34	2.86	3.55
824124		0.29	4.84	3.47	4.24	4.08	1.11	3.55
837953		2.95	2.00	4.04	3.97	5.08	2.63	3.55
511521		9.58	2.43	4.75	3.58	3.11	3.86	3.54
1493252		2.79	3.88	4.49	1.98	5.20	2.16	3.54
626640		2.35	2.29	3.09	5.39	3.53	3.42	3.54
2273		1.26	3.38	2.26	3.14	6.00	2.92	3.54
84023		1.97	5.71	3.40	2.86	3.42	2.30	3.54
84130		2.09	2.18	2.19	10.28	1.46	1.58	3.54
15026			3.31	4.40	3.92	2.96	3.05	3.53
95046			0.81	1.10	5.41	4.59	5.73	3.53
25101			2.89	2.88	6.72	1.14	4.00	3.52
6765			1.22	3.32	2.33	6.57	4.11	3.51
59240			0.83	2.74	2.70	8.19	3.04	3.50
75575			3.23	1.83	2.12	7.95	2.36	3.50
142079			2.46	3.91	3.17	3.47	4.47	3.49
84335			2.97	3.18	2.69	3.78	4.80	3.48
160434			1.56	2.15	1.34	9.00	3.33	3.48
25816	7 17.24	4.96	3.26	5.42	3.43	2.37	2.90	3.48

⊺3ble 1-1

308466	9.26	2.67	4.61	4.96	3.31	1.52	2.95	3.47
272576	6.22	1.80	1.34	3.85	4.11	4.71	3.25	3.45
108265	7.24	2.10	4.59	3.00	3.05	3.84	2.76	3.45
842784	326.54	94.88	0.85	3.11	1.95	7.01	4.30	3.44
1467207	3.68	1.07	2.33	4.23	4.08	2.71	3.86	3.44
784225	10.62	3.09	2.36	4.44	3.93	2.57	3.89	3.44
430179	51.95	15.13	2.52	2.04	9.04	1.80	1.78	3.44
796513	17.83	5.20	4.20	3.32	2.84	5.24	1.56	3.43
83363	5.08	1.48	3.97	3.58	1.64	3.63	4.33	3.43
489109	7.44	2.17	2.33	3.11	5.81	2.19	3.71	3.43
812977	7.62	2.22	2.38	2.15	6.48	4.79	1.35	3.43
1492238	13.96	4.08	4.19	3.72	2.61	4.20	2.41	3.43
740914	7.20	2.11	5. <b>02</b>	1.70	6.97	1.58	1.84	3.42
1475797	85.21	24.92	1.93	2.41	2.93	7.69	2.14	3.42
897042	4.95	1.45	6.12	3.62	2.70	3.57	1.07	3.42
249705	62.65	18.34	2.89	3.73	2.52	4.92	3.02	3.42
1476065	6.33	1.86	2.32	4.22	1.08	8.04	1.42	3.41
251731	0.82	0.24	2.88	3.28	3.37	3.75	3.78	3.41
838366		3.27	2.46	3.30	3.45	3.28	4.57	3.41
855799		2.64	2.70	3.06	2.67	2.66	5.96	3.41
701112		2.55	2.26	4.01	3.63	4.37	2.78	3.41
26406		1.92	2.41	3.58	3.68	3.80	3.58	3.41
727251		10.07	2.95	3.27	4.52	4.01	2.30	3.41
252453		5.91	2.32	4.20	2.91	5.06	2.54	3.41
1435185	1 1	0.75	5.55	2.80	3.10	3.49	2.08	3.40
741474		1.45	2.81	2.16	3.51	5.40	3.13	3.40
289891	1 1	2.30	3.60	2.40	7.51	1.93	1.56	3.40
796774	1 1	0.13	3.55	2.86	3.04	3.54	4.02	3.40
290642		2.28	4.70	3.05	2.60	3.37	3.27	3.40
14901		2.30	2.27	1.81	4.99	1.02	6.89	3.40
74529		8.59	7.43	1.72	1.39	4.51	1.94	3.40
95060		1.04	1.95	2.17	2.24	8.88	1.70	3.39
87840		2.13	3.07	4.92	1.88	5.02	2.03	3.38
95043		2.22	4.11	4.50	2.71	2.81	2.76	3.38
3982		1.73	2.98	3.80	3.61	3.36	3.14	3.38
84313	9 39.96	11.84	4.71	1.39	4.05	2.96	3.77	3.38
13718		1.56	2.28	4.02	3.12	4.62	2.80	3.37
29652	9.83	2.92	3.71	4.18	2.72	2.15	4.08	3.37
54463		0.84	4.23	3.65	3.42	1.95	3.59	3.37
84114	10 2.47	0.73	4.27	4.30	2.28	3.23	2.75	3.36
4977	28 1.27	0.38	4.06	3.75	2.82	2.96	3.23	3.36
16388	52 1.21		3.84	4.80	3.21	2.46	2.52	3.36
510			3 <b>.25</b>	1.40	2.40	2.89	6.87	3.36
7530			2. <b>28</b>	4.55	3.07	4.01	2.88	3.36
8982			2.34	3.18	3.05	4.51	3.70	3.36
4907			2.63	3.40	2.73	6.39	1.60	3.35
8556			1.02	1.19	0.83	1.13	12.53	3.34
7683			4.43	3.97	3.60	2.60	2.09	3.34
2121			5.13	3.22	4.38	1.81	2.15	3.34
3772			2.94	2.21	9.10	0.91	1.52	3.34
8098	76 27.7	4 8.32	4.70	3.46	1.68	2.47	4.37	3.34

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462325	8.76	2.63	2.80	0.87	2.51	7.67	2.81	3.33
624785	6.17	1.86	3.38	3.81	2.96	4.60	1.86	3.32
291345	70.40	21.23	7.87	1.97	3.76	1.35	1.64	3.32
257382	12.80	3.86	3.91	3.02	5.42	2.19	2.05	3.32
700900	9.94	3.00	3.26	2.88	3.29	3.74	3.40	3.31
742581	1.20	0.36	3.04	4.31	3.03	3.03	3.14	3.31
811867	24.57	7.43	3.72	3.16	2.52	4.66	2.46	3.31
502622	18.65	5.65	2.32	2.97	3.27	5.80	2.16	3.30
666371	0.85	0.26	3.76	2.72	3.71	3.35	2.97	3.30
362829	3.03	0.92	3.18	5.12	3.27	3.18	1.73	3.30
129146	7.97	2.42	3.78	3.62	2.54	2.98	3.55	3.30
135083	17.07	5.19	3.09	3.47	4.93	2.81	2.15	3.29
854678	6.21	1.89	1.76	9.64	1.93	1.21	1.90	3.29
321470	7.74	2.36	3.41	4.28	3.25	1.89	3.60	3.29
1472638	7.11	2.16	2,45	3.79	2.73	4.52	2.93	3.29
242952	16.94	5.16	3.84	3.11	3.39	3.54	2.54	3.28
884657	6.68	2.04	3.32	2.87	2.28	4.95	2.99	3.28
40338	2.61	0.80	4.02	2.85	3 <b>.77</b>	2.65	3.10	3.28
461670	8.90	2.71	2.03	4.16	4.85	2.37	2.99	3.28
45564	1.66	0.51	2.98	5.39	3.62	1.74	2.64	3.28
813637	34.08	10.41	3.65	3.22	2.20	5.54	1.75	3.27
291255	15.20	4.65	3.07	3.01	5.02	2.76	2.48	3.27
310493	7.56	2.32	3.61	3.65	3.06	2.77	3.24	3.26
433350	28.82	8.83	1.93	2.66	2.67	7.86	1.21	3.26
781454	9.75	2.99	2.79	1.30	9. <b>94</b>	1.22	1.06	3.26
685516	3.36	1.03	3.63	3.79	2.76	4.08	2.03	3.26
112785	7.75	2.38	2.21	3.83	3.58	3.82	2.81	3.25
264640	13.22	4.07	3.98	3.82	1.83	1.43	5.20	3.25
202904	12.77	3.93	2.44	4.34	2.65	3.34	3.48	3.25
251330	9.21	2.84	2.30	2.37	2.73	4.39	4.44	3.25
815164	0.88	0.27	3.76	3.41	4.07	3.68	1.32	3.25
1456602	6.11	1.88	2.40	2.64	2.44	4.72	4.03	3.25
1486260	13.23	4.08	2.75	2.87	3.17	5.07	2.36	3.24
322561	107.98	33.31	2.93	2.03	1. <b>79</b>	4.79	4.67	3.24
416436	11.08	3.42	5.36	3.73	2.74	3.15	1.23	3.24
827152	3.50	1.08	1.39	5.25	3.59	4.05	1.92	3.24
289818		2.31	2.49	4.60	3.59	3.30	2.19	3.23
273546		2.81	4.41	3.56	2.54	2.67	2.98	3.23
75886		4.37	5.08	3.68	1.09	4.18	2.11	3.23
416010		16.66	2.41	1.81	8.72	1.50	1.69	3.23
838628		0.88	2.12	3.44	3.70	3.20	3.67	3.23
22747			2.96	3.25	2.59	4.14	3.15	3.22
951091		0.63	2.71	3.17	4.10	2.80	3.30	3.22
35812			2.94	2.49	2.99	3.34	4.32	3.22
82874			4.36	3.56	2.21	3.84	2.10	3.21
1631713			2.81	3.64	3.90	2.77	2.93	3.21
263070			4.55	4.07	1.99	3.33	2.10	3.21
1031698			2.99	2.12	2.54	4.81	3.58	3.21
1602120			1.98	3.03	2.64	5.10	3.30	3.21
454672			5.39	3.57	1.75	2.15	3.16	3.20
837904	103.13	32.18	2.10	2.12	2.01	4.30	5.50	3.20

Table 1-1

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34869	2.74	0.85	1.85	2.96	1.76	7.56	1.87	3,20
840590	21.70	6.78	2.44	1.18	2.39	8.32	1.68	3.20
377672	2.59	0.81	0.93	0.98	0.68	12.46	0.96	3.20
511632	7.85	2.46	2.42	2.96	2.15	4.52	3.94	3.20
531028	21.14	6.62	3.62	3.65	3.05	2.41	3.26	3.20
321386	2.45	0.77	3.26	2.66	2.50	4.45	3.11	3.20
754221	6.80	2.13	2.49	3.95	5.19	2.42	1.93	3.20
433573	10.67	3.34	3.00	3.46	3.80	2.33	3.38	3.19
489823	33.49	10.49	2.75	4.23	1.58	5.36	2.05	3.19
248454	3.70	1.16	7.24	2.69	2.10	2.69	1.24	3.19
298155	2.85	0.89	2.52	4.31	2.59	3.88	2.66	3.19
530814	46.55	14.60	0.83	4.74	4.23	4.56	1.60	3.19
294259	4.21	1.32	3. <b>87</b>	2.34	3.12	1.09	5.52	3.19
726488	10.10	3.17	3.31	3.38	3.27	3.50	2.45	3.18
511459	37.04	11.64	2.30	3.42	4.76	3.38	2.06	3.18
503468	51.40	16.15	3.74	2.80	1.75	4.34	3.28	3.18
756600	47.06	14.79	3.37	1.74	1.79	4.21	4.80	3.18
760344	5.89	1.85	3.62	4.43	2.41	1.42	4.03	3.18
753252	7.78	2.45	2.52	2.78	2.96	4.05	3.59	3.18
26230	5.08	1.60	1.18	3.71	4.40	3.50	3.09	3.18
37980	0.84	0.27	1.41	2.73	1.73	8.06	1.95	3.18
156386	2.14	0.67	3.02	3.99	3.91	2.87	2.09	3.18
969844	9.12	2.88	5.15	2.28	2.19	3.92	2.33	3.17
855723	6.73	2.12	2.86	3.79	2.80	3.00	3.38	3.17
1240220	7.60	2.40	0.79	2.71	6.02	4.99	1.31	3.17
451769	5.25	1.66	3.67	2.29	2.52	5.57	1.78	3.16
810027	0.44	0.14	3.75	3.05	2.90	3.82	2.31	3.16
731272	5.02	1.59	2.45	2.89	3.95	4.89	1.62	3.16
129024	4.30	1.36	5.53	0.58	1.77	1.86	6.04	3.16
795907	5.39	1.71	1.43	1.64	3.37	0.61	8.72	3.15
731257	4.45	1.41	1.78	2.17	3.22	7.14	1.45 3.63	3.15
703707 811603		6.22	2.6 <b>8</b> 3. <b>87</b>	2.65 2.99	3.51 2.25	3.30 3.41	3.03	3.15 3.15
884346		1.71	2.20	4.78	2.25 3.45	3.41	2.15	3.15
201628		2.39	4.15	2.46	3.43 1.74	5.00	2.35	3.14
840821		5.60	3.31	3.10	3.72	2.88	2.71	3.14
344432		4.93	3.94	2.68	2.53	2.44	4.11	3.14
590853		0.21	2.61	2.78	3.43	3.40	3.47	3.14
767298		6.18	2.43	3.33	2.32	4.12	3.48	3.14
36387		1.02	2.66	3.88	4.50	1.19	3.46	3.14
302955		3.39	1.90	5.11	2.20	4.12	2.36	3.14
796885		3.08	4.15	1.95	2.26	4.54	2.77	3.13
725489		8.54	3.93	3.59	2.81	3.60	1.72	3.13
121489			3.79	3.88	3.08	2.54	2.35	3.13
263894			3.70	3.08	2.13	4.71	2.03	3.13
782383	1		2.40	2.83	2.35	5.64	2.42	3.13
815563			3.37	3.78	2.43	4.15	1.90	3.13
245547			1.15	5.91	3.40	3.50	1.66	3.12
85657			2.42	1.67	5.16	3.07	3.31	3.12
25175			2.40	2.73	2.75	5.27	2.45	3.12
35699			4.19	2.74	2.51	3.63	2.54	3.12
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Table 1-1

502463	5.77	1.85	2.78	3.60	3.10	2.63	3.49	3.12
1461528	3.95	1.27	3 <b>.05</b>	2.21	5.36	2.64	2.33	3.12
244652	24.70	7.92	2.62	1.81	7.76	1.54	1.87	3.12
266106	20.68	6.63	2.76	3.21	6.30	1.40	1.93	3.12
1415981	4.13	1.32	3.58	3.02	4.59	2.33	2.07	3.12
66902	3.21	1.03	4.24	2.05	1.56	0.87	6.87	3.12
283919	7.56	2.43	7.31	3.63	1.47	2.02	1.15	3.11
813712	7.62	2.45	1.14	3.64	1.91	6.70	2.18	3.11
195903	6.95	2.24	2.45	4.56	3.51	2.97	2.05	3.11
345032	20.78	6.69	1.19	4.73	3.52	2.65	3.46	3.11
1635665	5.13	1.65	4.57	3.12	0.90	5.26	1.69	3.11
154749	1.76	0.57	3.34	3.27	3.77	2.57	2.58	3.11
287728	2.76	0.89	0.59	4.01	0.98	9.25	0.70	3.11
701409	1.54	0.50	4.21	2.63	3.95	2.39	2.34	3.10
627687	56.19	18.11	1.06	1.86	3.60	4.82	4.16	3.10
855244	15.78	5.09	6.04	0.82	7.22	0.56	0.88	3.10
293243	13.17	4.25	4.44	2.03	6.12	0.95	1.96	3.10
295206	3.16	1.02	5.72	3.11	1.43	2.48	2.76	3.10
744846	2.69	0.87	3.69	4.51	2.68	3.60	1.01	3.10
251645	0.50	0.16	3.12	3.21	3.74	3.30	2.12	3.10
357364	0.72	0.23	5.76	3.07	1.81	3.15	1.68	3.09
451707	7.59	2.46	2.46	1.83	6.35	1.34	3.48	3.09
250673	2.66	0.86	7.23	1.34	2.15	1.85	2.88	3.09
1031086	25.15	8.15	3.21	2.15	2.25	5.00	2.83	3.09
50892	7.95	2.58	3.62	2.53	3.85	2.69	2.71	3.08
52327	5.70	1.85	1.94	3.44	2.70	3.46	3.87	3.08
448930	7.53	2.45	2.47	2.25	2.41	3.62	4.66	3.08
713109	2.44	0.79	1.97	2.95	3.60	4.59	2.28	3.08
815794	14.69	4.77	5.65	1.93	1.67	1.95	4.18	3.08
7890 <b>9</b> 1	9.50	3.09	6.20	4.45	1.91	1.88	0.95	3.08
950836	34.79	11.31	2.78	1.53	8.20	0.92	1.95	3.08
213890	14.14	4.60	2.37	5.07	2.41	3.28	2.24	3.07
74223	5.42	1.76	3.29	3.47	3.41	2.68	2.52	3.07
41174		1.31	3.25	3.60	2.82	3.70	1.98	3.07
1635874	6.06	1.97	6.28	3.19	1.83	2.38	1.69	3.07
810284		0.74	5.17	2.15	2.18	2.13	3.71	3.07
845502		1.26	1.42	3.27	3.03	4.57	3.06	3.07
461489		0.30	3.19	3.40	2.48	2.98	3.28	3.07
361097		1.84	3.06	3.81	1.75	3.06	3.64	3.07
139113		1.41	2.39	3.80	2.88	5.14	1.10	3.06
813636		27.59	3.51	2.53	2.58	4.58	2.11	3.06
743804		1.66	3.74	2.59	2.21	3.39	3.38	3.06
486567		2.99	2.48	4.29	2.98	2.50	3.06	3.06
665127		1.79	3.90	3.70	2.31	2.65	2.73	3.06
136605			3.42	0.62	2.86	2.82	5.58	3.06
395625		4.73	2.38	3.22	3.22	3.25	3.22	3.06
287528		0.65	3.56	0.96	5.02	3.66	2.06	3.06
323 <b>47</b> 4 884388		0.41 1.60	5.34	3.25	2.05	2.36	2.27	3.06
22252			2.97	2.78	2.75	3.38	3.40	3.06
260628			1.68	4.03	3.69	4.09	1.79	3.05
200020	PI 6.74	1 2.73	1.68	4.70	3.88	2.83	2.17	3.05

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Table 1-1

471266	2.72	0.89	3.15	2.95	2.66	3.45	3.03	3.05
565062	1.38	0.45	1.72	3.15	3.97	2.87	3.53	3.05
701532	3.26	1.07	3.06	2.86	3.51	1.98	3.84	3.05
878127	3.53	1.16	2.07	3.35	4.53	3.32	1.96	3.05
878373	31.06	10.19	3.07	3.84	2.03	4.00	2.29	3.05
40075	3.13	1.03	2.09	3.44	2.70	5.00	2.00	3.05
884766	9.95	3.27	1.22	3.47	2.45	2.67	5.40	3.04
754192	3.30	1.09	4.21	2.70	2.65	3.45	2.21	3.04
299154	1.37	0.45	1.64	3.95	3.01	2.17	4.42	3.04
344589	53.28	17.55	1.28	4.11	3.84	2.63	3.31	3.04
897153	10.77	3.55	3.27	3.89	4.14	2.12	1.76	3.04
328613	0.46	0.15	2.60	3.52	3.31	2.33	3.42	3.03
884867	24.71	8.14	3.57	3.43	1.95	3.39	2.83	3.03
289288	5.44	1.79	3.47	2.18	3 <b>.94</b>	1.96	3.62	3.03
788518	13.97	4.61	2.21	4.58	2.41	3.19	2.78	3.03
825842	6.00	1.98	1.78	2.66	4.19	1.45	5.08	3.03
133130	11.12	3.67	3.58	0.72	2.94	2.70	5.20	3.03
252864	1.20	0.40	2.46	3.75	2.39	3.50	3.03	3.03
813187	22.70	7.51	1.31	2.52	4.98	3.97	2.34	3.02
1635110	38.13	12.62	4.50	2.40	3.02	2.46	2.73	3.02
266747	3.05	1.01	3.26	3.57	2.70	0.99	4.59	3.02
490360	5.94	1.97	2.87	3.33	2.69	3.50	2.72	3.02
951216	2.02	0.67	3.88	2.20	2.08	4.19	2.75	3.02
322914	9.34	3.10	4.67	3.02	2.07	1.93	3.39	3.02
845609	15.29	5.07	3.85	3.88	2.84	3.05	1.46	3.02
1467108	4.29	1.42	1.21	3.61	4.50	3.23	2.53	3.02
415978	24.58	8.16	2.76	1.42	3.19	4.93	2.76	3.01
645368	8.48	2.82	2.47	3.08	3.32	2.42	3.76	3.01
854899	18.74	6.24	3.77	2.48	2.61	2.49	3.67	3.00
80764	26.16	8.72	2.38	2.68	2.24	3.93	3.77	3.00
1501998	2.03	0.68	3.20	3.50	2.70	3.57	2.05	3.00
590298	10.43	3.47	3.90	3.96	3.21	2.04	1.90	3.00
594693	7.33	2.44	3.84	2.11	7.17	1.01	0.88	3.00
856115	1.30	0.43	9.13	1.80	2.06	0.91	1.10	3.00
364022 730412	4.15 1.25	1.39 0.42	3.12	5.09	2.52	2.61	1.64	3.00
843140	14.94	4.99	2.86	2.00	3.95 3.94	3.63	2.55	3.00
342211	1.92	0.64	1.67 2. <b>7</b> 5	3.33 3.24	2.28	2.97	3.06 1.90	3.00 3.00
742806		0.59	2.73	3.24	2.26 4.16	4.81 2.33	2.85	2.99
884480		3.91	3.07	2.74	1.85	4.27	3.01	2.99
1636707		6.44	2.58	3.23	2.07	3.98	3.07	2.99
712610			3.02	3.36	3.85	3.46	1.23	2.98
360232			1.98	3.91	3.63	3.66	1.72	2.98
74511			4.10	3.44	2.52	0.97	3.89	2.98
343167		5.24	3.15	4.46	2.96	2.05	2.27	2.98
39885	•		0.88	2.80	4.66	4.76	1.80	2.98
122183			2.66	4.22	2.01	2.53	3.46	2.98
38253			3.19	4.23	2.69	2.79	1.98	2.98
745103			2.03	3.44	3.39	2.05	3.96	2.97
155838			2.70	1.94	5.31	2.18	2.74	2.97
813149			5.57	3.70	2.01	2.38	1.21	2.97
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Table 1-1

469685	4.95	1.67	3.25	3.34	2.23	3.52	2.51	2.97
824640	6.96	2.35	1.37	1.54	3.69	3.65	4.59	2.97
825833	1.95	0.66	3.29	2.75	4.19	2.65	1.97	2.97
25517	8.52	2.88	3.15	3.29	2.67	2.43	3.28	2.97
842767	2.15	0.73	2.59	3.85	3.19	2.67	2.53	2.96
148225	3.89	1.31	2.29	2.66	1.88	5.50	2.47	2.96
772304	113.14	38.24	3.78	2.19	2.18	2.83	3.83	2.96
246800	6.67	2.26	2.06	3.22	2.26	4.33	2.91	2.96
322192	7.97	2.69	1.72	3.28	5.02	1.74	3.02	2.96
755765	4.01	1.36	2.43	3.91	4.37	1.46	2.62	2.96
462468	3.24	1.10	3.23	3.74	3.40	1.72	2.69	2.96
701402	4.23	1.43	1.82	3.11	5.63	2.91	1.31	2.96
253253	1.73	0.59	4.71	4.49	1.70	2.17	1.68	2.95
211206	4.71	1.60	2.58	3.21	2.52	2.63	3.82	2.95
813825	0.52	0.18	3.85	3.04	1.69	4.84	1.32	2.95
898227	2.02	0.69	1.45	4.11	2.91	4.09	2.18	2.95
45578	5.05	1.72	4.61	2.16	2.26	3.42	2.27	2.94
1522799	1.23	0.42	2.25	3.40	2.72	3.47	2.86	2.94
593520	6.25	2.13	2.05	2.67	3.01	4.62	2.34	2.94
487773	29.51	10.06	4.78	2.56	4.05	1.34	1.95	2.93
502690	4.09	1.39	2.22	3.28	2.51	4.37	2.29	2.93
731104	0.65	0.22	3.42	2.35	3.12	2.70	3.07	2.93
796341	7.16	2.44	2.21	2.93	2.48	3.55	3.49	2.93
1492304	4.23	1.44	2.24	2.81	3.19	4.05	2.36	2.93
77805	15.55	5.31	2.32	4.51	2.41	2.27	3.13	2.93
415700	3.36	1.15	2.29	2.89	4.79	1.53	3.14	2.93
701496	2.28	0.78	2.43	2.22	5.16	3.20	1.62	2.93
233419	5.55	1.90	1.88	3.96	3.98	1.76	3.06	2.93
767994	4.29	1.47	2.76	1.52	8.75	0.69	0.92	2.93
254310	10.26	3.51	3.10	2.50	6.68	1.35	1.00	2.92
48033	0.51	0.18	2.77	3.62	3.02	2.44	2.76	2.92
811842		3.39	2.78	3.26	2.72	2.81	3.05	2.92
44975	1 1	3.05	2.47	4.15	2.22	2.69	3.08	2.92
243068		3.20	2.27	2.18	2.15	5.77	2.23	2.92
1635350		3.48	2.60	2.63	5.83	2.20	1.33	2.92
759948	1 1	1.39	2.12	1.92	1.94	7.00	1.59	2.91
35807		1.00	3.30	2.96	2.26	3.58	2.45	2.91
45515		1.33 16.97	1.58	3.77	5.25	1.91	2.04	2.91
1591 <b>62</b> 2 460 <b>84</b> 1		0.35	2.27 4.19	3.52 1.78	1.93 <b>4.04</b>	4.64 2.42	2.17	2.91 2.91
1411726		10.55	3.69	2.02	2.19	4.86	2.11 1.78	2.91
160485	1 1	1.78	3.02	1.25	1.91	2.64	5.72	2.91
74269		0.63	2.56	3.08	2.90	4.14	1.83	2.90
242700	1 .	1.78	3.05	3.96	1.69	3.81	1.99	2.90
868838			1.07	1.24	0.96	9.00	2.21	2.90
142083			1.37	2.47	2.52	4.60	3.53	2.90
82577		4.19	2.16	5.30	2.82	2.12	2.09	2.90
82610			2.02	1.86	7.90	1.13	1.56	2.89
46148	4		3.29	3.33	2.51	2.75	2.59	2.89
4948			3.81	2.49	2.73	3.24	2.15	2.89
12972			2.93	1.99	4.17	3.16	2.17	2.89
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Table 1-1

627273	1.58	0.55	1.86	3.85	2.31	3.67	2.73	2.88
143306	10.42	3.62	3.88	2.52	1.98	3.46	2.55	2.88
730313	3.95	1.37	2.62	2.27	3.50	2.08	3.93	2.88
825718	4.03	1.40	2.44	2.50	3 <b>.7</b> 7	3.02	2.67	2.88
1591264	3.18	1.11	4.68	4.17	2.19	2.11	1.23	2.88
784016	33.80	11.75	2.08	4.19	3.69	2.77	1.66	2.88
41940	10.51	3.65	1.79	2.81	3.34	3.64	2.80	2.88
841501	5.94	2.07	3.74	4.22	2.48	2.62	1.31	2.88
853968	12.10	4.21	3.42	3.85	5.08	1.19	0.84	2.87
489663	2.11	0.73	3.88	2.51	1.67	3.15	3.17	2.87
669371	4.99	1.74	2.07	1.98	7.30	1.15	1.87	2.87
815760	2.31	0.81	3.18	2.00	4.13	3.96	1.09	2.87
125722	3.28	1.14	4.52	2.27	2.75	3.95	0.86	2.87
897625	2.58	0.90	1.23	2.12	4.08	3.14	3.79	2.87
768179	3.80	1.32	1.92	3.06	2.43	4.11	2.83	2.87
768469	4.17	1.45	2.78	1.73	2.41	3.64	3.79	2.87
898265	20.16	7.03	1.31	2.31	3.96	2.43	4.34	2.87
868119	10.00	3.49	2.81	3.31	3.66	1.82	2.73	2.87
897983	26.99	9.42	2.98	3.14	2.24	2.33	3.64	2.86
1416086	8.24	2.88	3.59	2.09	5.52	1.57	1.54	2.86
141972	18.88	6.60	3.58	3.00	3.07	1.99	2.68	2.86
144977	71.81	25.09	2.67	3.73	3.36	1.42	3.14	2.86
949940	12.52	4.38	4.69	2.75	1.41	3.20	2.25	2.86
898251	2.02	0.71	2.79	2.93	3.12	4.01	1.45	2.86
80500	140.81	49.33	0.91	2.65	1.64	5.24	3.85	2.86
823588	7.72	2.70	2.44	3.65	3.22	2.51	2.45	2.85
269300	10.64	3.73	1.47	2.55	4.46	4.07	1.71	2.85
823627	3.25	1.14	2.72	3.08	3.03	1.90	3.53	2.85
1589973	5.17	1.81	2.56	3.21	1.99	4.51	1.99	2.85
246541	10.77	3.78	2.76	3.03	2.41	3.92	2.14	2.85
784253	11.76	4.12	2.36	3 <b>.6</b> 7	2.76	2.44	3.04	2.85
27072	0.29	0.10	2.61	2.23	3.44	2.93	3.04	2.85
754449	1.78	0.63	2.10	3.63	2.52	4.00	1.99	2.85
82905	1.21	0.43	3.27	1.88	1.90	3.49	3.70	2,85
743828	3.34	1.17	1.16	1.69	5.22	0.99	5.18	2.85
743851	8.22	2.89	3.40	2.79	2.54	3.77	1.72	2.84
1475746	12.48	4.39	3.24	2.13	2.29	3.86	2.70	2.84
283196		1.34	1.51	2.93	1.19	6.19	2.38	2.84
866276		0.68	1.99	3.31	2.96	2.62	3.33	2.84
144926		3.24	3.79	2.02	2.76	3.87	1.75	2.84
769926			6.19	1.95	2.54	2.39	1.09	2.83
730554			2.36	3.04	1.33	4.78	2.64	2.83
40017			2.97	3.32	1.84	3.86	2.12	2.82
897577			2.18	2.74	3.06	3.81	2.31	2.82
1492412	B.	1 1	1.73	2.18	3.44	2.84	3.92	2.82
1336262	1		4.41	2.34	4.62	1.32	1.41	2.82
250678			3.14	3.84	3.52	1.72	1.86	2.82
810220			4.33	3.00	1.34	3.50	1.92	2.82
271076		: 1	2.34	3.42	3.53	2.75	2.04	2.82
111812			3.30	4.08	4.25	1.15	1.29	2.81
898 <b>09</b> 6	4.06	1.45	3.41	2.02	3.72	1.08	3.83	2.81

Table 1-1

250869	123.92	44.10	8.11	2.05	1.68	0.76	1.46	2.81
840620	13.81	4.92	1.46	1.93	2.95	3.21	4.47	2.81
53165	0.89	0.32	3.53	3.16	2.83	2.33	2.16	2.80
784128	1.83	0.65	1.61	1.75	6.85	1.95	1.84	2.80
452668	4.95	1.77	2.13	1.80	2.80	5.84	1.44	2.80
811911	11.89	4.25	3.15	1.99	3.02	2.42	3.41	2.80
854122	4.20	1.50	5.92	3.38	2.72	0.79	1.18	2.80
491184	6.35	2.27	3.28	1.15	8.07	0.69	0.80	2.80
825356	1.18	0.42	2.48	2.51	4.19	3.57	1.22	2.79
214884	3.55	1.27	4.42	2.54	1.68	3.83	1.50	2.79
450711	1.29	0.46	2.21	3.64	2.22	4.08	1.82	2.79
1593502	2.25	0.81	1.62	3.22	2.57	2.01	4.54	2.79
825058	3.42	1.23	2.71	1.17	8.02	1.37	0.68	2.79
285226	25.74	9.23	7.63	1.20	2.23	1.77	1.12	2.79
191904	7.67	2.75	3.12	2.55	2.70	2.27	3.28	2.79
786609	4.81	1.73	0.70	0.71	2.56	9.16	0.80	2.78
204146	2.40	0.86	1.54	5.79	2.84	1.41	2.34	2.78
713286	0.89	0.32	3.32	2.65	3.34	3.52	1.08	2.78
416833	6.07	2.18	1.35	3.81	1.74	4.90	2.09	2.78
268211	3.43	1.23	2.33	1.79	1.77	1.26	6.74	2.78
782439	35.89	12.92	3.09	2.38	1.70	3.98	2.74	2.78
810741	72.43	26.10	3.64	2.64	2.20	3.00	2.38	2.78
431286	13.81	4.98	2.22	2.98	3.24	1.94	3.50	2.77
826335	2.24	0.81	2.09	2.46	3.66	3.16	2.49	2.77
796313	0.78	0.28	3.22	2.69	2.55	2.13	3.26	2.77
812989	2.18	0.79	2.58	2.17	3.23	3.95	1.93	2.77
258521	3.33	1.20	1.48	7.97	1.12	1.13	2.14	2.77
526945		0.41	3.17	1.74	2.47	3.11	3.36	2.77
197821	5.95	2.15	5.83	4.39	1.04	1.29	1.28	2.77
1631747		1.84	4.14	4.87	1.99	1.76	1.08	2.77
1475738		86.12	3.33	2.08	2.23	3.93	2.25	2.76
248261		0.55	4.72	1.35	0.99	1.14	5.61	2.76
841348		4.33	4.50	2.03	0.87	4.69	1.72	2.76
1031346		1.16	3.61	2.85	2.26	3.16	1.91	2.76
244154		1.79	2.49	2.18	3.84	2.06	3.22	2.76
245860		11.02	5.36	1.24	5.40	0.69	1.09	2.76
271472		4.28	2.07	3.57	2.66	2.40	3.07	2.76
753917		2.36	2.95	3.41	1.98	3.47	1.95	2.75
51749			2.28	1.01	1.83	2.34	6.28	2.75
296429			1.94	3.44	2.46	2.47	3.43	2.75
430318			1.77	2.79	3.60	1.97	3.59	2.74
283142			3.20	3.03	3.74	1.95	1.78	2.74
43534			3.20	2.13	2.58	3.61	2.19	2.74
27853			1.66	3.41	1.66	4.44	2.52	2.74
48972			2.33	2.25	3.98	3.06	2.07	2.74
81494			2.62	3.11	2.36	3.47	2.14	2.74
50374			1.17	3.06	1.97	5.66	1.82	2.74
163214	1		2.34	3.20	3.06	2.96	2.13	2.74
42821			3.87	3.41	1.92	2.28	2.19	2.74
87831			2.19	2.60	1.52	4.16	3.20	2.73
157308	7 2.41	0.88	1.63	2.09	4.06	4.16	1.72	2.73

Table 1-1

841399	9.08	2 221	2.44	2.07	2.40	2 2 1	2.45	2.72
897730	41.12	3.33 15.07	2.44	2.07 1.82	3.48 2.32	3.21 2.54	3.88	2.73
866882	10.73	3.94	3.07 1.81	4.01	2.32 3.71	1.83	2.28	2.73 2.73
1160558	8.03	2.94		3.29	2.53	3.66	2.00	2.73
127928	3.67	1.35	2.15		2.33 2.89	2.15	3.51	2.73
	11.55		1.79	3.28 2.82	3.44	2.74	3.21	2.73
897971		4.24	1.43			4.48	2.08	
447568	9.00	3.31	3.49	1.70	1.86		1.30	2.72
811880	3.13	1.15	1.85	2.58	4.75	3.12	1.85	2.72
810725	3.27	1.20	1.92	2.23	1.63	5.97	2.38	2.72
197176	20.79	7.65	3.06	2.51	3.48	2.16	1	2.72 2.72
758347	. 2.75	1.01	1.61	3.59	3.19	2.80	2.41	
43705	0.35	0.13	2.72	3.74	1.90	3.12		2.71
813280	4.14	1.53	1.72	2.58	2.29	3.06	3.91	2.71
195652	18.03	6.65	2.18	3.01	2.07	4.00	2.29	2.71
868308	65.99	24.35	3.48	1.83	1.37	4.88	2.00	2.71
429906	5.35	1.98	1.98	3.10	2.51	2.18	3.77 3.64	2.71
83011	32.97	12.18	2.39	2.09	2.06	3.35		2.71
838611	52.82	19.51	4.00	0.88	1.23	4.84	2.60	2.71
783697	3.19	1.18	2.47	3.13	3.04	2.62	2.26	2.70
268178	28.02	10.37	1.92	3.23	2.46	3.05	2.85 6.21	2.70
460487	24.08	8.92	0.68	0.85	5.68	0.08 3.44	2.70	2.70 2.70
418053	7.57	2.80	3.25	1.89	2.22	3.4 <del>4</del> 3.14	2.68	2.70
884894	41.65	15.44	3.11	2.60	1.96	5.02	1.74	2.70
824066	4.08	1.51	2.48	2.33	1.91	3.54	2.39	2.70
950450		0.66	1.96	2.45	3.15	3.3 <del>4</del> 9.45	1.06	2.70
50344		0.32 2.12	1.00 2.14	1.06	0.91 2.38	2.44	3.04	2.69
42076			2.14	3.48	2.36	3.37	1.62	2.69
68636		1.80 0.43	1.89	3.27 2.57	2.43 3.3 <b>8</b>	3.81	1.78	2.69
26185 278687	1 1	4.88	2.06	4.03	2.38	1.89	3.07	2.69
391984	1	1.70	3.22	3.44	1.67	2.03	3.06	2.68
279785	1 1	0.34	1.97	1.94	1.70	5.89	1.92	2.68
8971 <b>9</b> 9		5.51	3.48	2.55	2.61	3.07	1.71	2.68
263697		1.58	3.06	3.23	4.13	1.12	1.87	2.68
770983		2.70	1.30	5.12	2.60	2.36	2.03	2.68
505344		0.84	3.69	2.24	1.96	3.13	2.38	2.68
25386		0.63	5.25	3.14	1.49	1.55	1.93	2.67
75269		2.27	3.98	2.14	0.98	4.23	2.02	2.67
85768		20.36	3.42	2.88	1.57	3.42	2.07	2.67
104903		1.15	2.50	3.09	3.57	3.21	0.99	2.67
74304		2.43	3.55	3.05	3.00	2.00	1.76	2.67
46176		3.24	2.63	1.55	4.32	3.11	1.72	2.67
5192		1.68	5.04	1.90	2.84	1.30	2.26	2.67
147182		37.33	2.97	2.21	1.62	5.24	1.30	2.67
30788			3.13	1.77	2.17	4.14	2.12	2.67
74308			4.46	3.27	1.20	2.05	2.35	2.67
25145	•		1.29	2.18	2.99	. 5.63	1.24	2.67
79573			3.12	3.28	1.60	3.79	1.52	2.66
83847			4.43	1.79	2.84	1.08	3.16	2.66
36132				1.41	2.10	6.37	1.76	2.66
52986	5.75	2.16	2.44	2.09	3.05	3.43	2.29	2.66

Table 1-1

486175	10.39	3.91	1.74	7.19	2.53	0.92	0.91	2.66
307029	56.38	21.22	3.05	1.98	1.59	4.58	2.09	2.66
243321	2.79	1.05	2.81	1.53	1.43	1.74	5.74	2.65
590398	12.27	4.63	4.11	2 <u>.2</u> 2	2.03	3.29	1.58	2.65
1475028	249.73	94.34	2.17	1.81	1.81	4.31	3.14	2.65
595604	9.41	3.56	1.72	1.96	4.39	3.06	2.10	2.65
188151	3.25	1.23	1.76	5.51	1.64	2.21	2.09	2.64
859832	2.57	0.97	1.88	3.53	2.79	3.19	1.81	2.64
42313	1.33	0.51	3.37	2.77	2.16	3.35	1.55	2.64
282144	30.16	11.43	1.47	2.73	2.45	5.25	1.30	2.64
32887	3.04	1.16	1.78	3.54	3.59	2.10	2.17	2.64
137254	1.35	0.51	3.82	2.00	4.43	1.38	1.55	2.64
234036	4.77	1.81	1.87	3.76	2.13	3.14	2.28	2.64
206785	6.05	2.30	4.91	3.70	1.64	1.42	1.50	2.63
244202	9.25	3.51	1.51	2.29	3.36	1.92	4.09	2.63
827144		0.96	1.88	3.18	2.21	1.64	4.24	2.63
838504		15.95	1.67	3.37	2.82	3.53	1.77	2.63
376764	2.58	0.98	1.55	2.16	3.58	3.03	3.03	2.63
744933	2.15	0.82	0.98	1.12	0.87	8.87	1.30	2.63
161456		5.08	0.92	0.15	10.09	0.13	1.83	2.63
564846		0.72	3.47	1.34	2.53	2.17	3.62	2.63
815072	3.49	1.33	5.03	2.36	1.34	2.07	2.32	2.62
1599489	5.51	2.10	3.32	3.51	1.94	2.95	1.40	2.62
80292	2.18	0.83	2.30	3.64	1.47	3.07	2.62	2.62
262865	30.48	11.64	3.50	2.06	4.94	0.99	1.61	2.62
590390	6.42	2.45	4.07	2.94	1.83	3.34	0.92	2.62
610011	34.63	13.23	3.40	2.31	2.21	1.95	3.23	2.62
855487	7 8.56	3.27	2.21	3.52	3.16	1.26	2.94	2.62
503091	7.27	2.78	1.87	1.80	7.01	1.54	0.86	2.62
825719		0.29	3.01	3.07	3.22	2.78	0.99	2.62
28272		3.18	3.44	1.29	2.89	2.24	3.20	2.61
76779		2.18	2.43	1.90	2.16	3.46	3.13	2.61
78412		2.50	1.21	2.18	5.46	2.80	1.41	2.61
23433	1 1	1.21	2.32	2.24	3.25	2.18	3.04	2.61
73897		6.13	1.21	2.29	7.41	1.31	0.82	2.61
4141		43.25	3.28	2.76	1.54	3.09	2.35	2.60
78426			1.81	3.29	2.15	3.61	2.13	2.60
81324			3.66	1.96	4.73	1.34	1.29	2.60
84053	1		3.14	3.14	1.61	3.33	1.76	2.60 2.59
81175			3.15	2.92	3.96	1.62	1.32	2.59
50236			3.58	2.45	2.17 2.12	3.01 2.49	1.74 3.44	2.59
74494			3.76	1.13			5.42	2.59
85563			1.55	1.65	3.18	1.13 2.94	1.65	2.58
7261			3.07	3.09	2.14		1.49	2.58
43610			2.43	2.14	3.01	3.83	3.67	2.58
4873			2.52	2.05	1.52	3.12	5. <b>8</b> 7	2.57
8141:	1	l I	1.83	1.77	1.85	1.52 5.65	1.19	2.57
4353		1 .	1.68 1.53	2.24 3.21	2.09 1.33	4.15	2.62	2.57
8385 1376			3.16	2.54	1.89	2.17	3.08	2.57
412				2.29	3.16	3.58	2.07	2.57
412	UT 2.03	ا0.00 اد	1.72	2.27	3.10	ەد.د	2.07	2.51

Table 1-1

				ţ.				
1558642	3.11	1.21	3.27	2.26	1.72	4.08	1.50	2.56
432194	10.34	4.04	1.58	3.07	2.91	1.91	3.34	2.56
232628	4.02	1.58	4.00	1.88	4.50	1.05	1.32	2.55
951022	3.02	1.18	3.66	2.02	3.80	1.61	1.67	2.55
809824	2.66	1.05	1.68	2.11	2.41	3.25	3.24	2.54
897906	10.34	4.08	1.96	2.02	4.28	0.64	3.78	2.54
809722	50.59	19.96	0.71	0.97	3.43	3.59	3.97	2.54
138865	5.42	2.14	1.16	3.20	2.53	3.12	2.65	2.53
825012	4.46	1.76	3.39	2.40	2.05	3.22	1.61	2.53
1472689	5.15	2.03	1.97	0.59	1.72	7.13	1.27	2.53
322537	3.18	1.26	1.26	6.07	2.04	0.89	2.40	2.53
291974	57.24	22.63	3.29	1.87	1.09	4.48	1.91	2.53
307231	2.04	0.81	1.95	1.61	1.41	6.48	1.19	2.53
1638479	3.27	1.30	2.08	3.88	3.23	1.78	1.67	2.53
415415	9.68	3.83	3.25	2.43	1.49	3.21	2.26	2.53
251417	0.69	0.27	3.59	3.13	1.67	1.79	2.46	2.52
452936	5.15	2.04	3.47	2.84	3.54	1.65	1.11	2.52
488301	0.97	0.38	2.10	1.37	1.81	5 <b>.57</b>	1.76	2.52
430235	3.43	1.36	4.46	3.50	1.54	1.77	1.34	2.52
784085	3.93	1.56	3.14	2.04	3.39	2.55	1.49	2.52
741988	1.74	0.69	1.90	3.05	2.20	3.46	1.99	2.52
627688	7.37	2.93	1.52	1.61	6.90	1.77	0.79	2.52
825859	1.64	0.65	1.30	2.26	2.23	3.48	3.31	2.52
884892	8.02	3.19	1.99	2.90	3.01	3.04	1.63	2.52
277660	14.16	5.65	1.86	2.15	2.18	3.13	3.22	2.51
76605	2.48	0.99	4.39	3.21	1.27	2.11	1.53	2.50
220069	4.81	1.92	1.69	3.29	3.04	1.86	2.61	2.50
460 <b>798</b>	1.30	0.52	1.95	1.79	3.77	1.74	3.24	2.50
247546	11.16	4.48	1.50	1.33	1.37	2.54	5.72	2.49
814080	8.19	3.29	1.76	1.45	2.40	1.52	5.32	2.49
1422794	2.62	1.05	0.95	1.99	3.82	3.31	2.38	2.49
51981	19.62	7.89	3.06	1.82	1.65	2.42	3.50	2.49
325606		2.93	3.10	2.79	1.65	3.28	1.62	2.49
277596		1.17	1.17	2.54	1.50	6.20	1.01	2.49
788205		5.32	3.46	2.10	1.43	1.78	3.65	2.48
796527		0.55	2.22	1.81	1.52	5.45	1.39	2.48
827132		2.68	2.47	1.83	1.62	3.44	3.03	2.48
842863		4.49	3.25	2.37	1.47	2.24	3.06	2.48
29964	1 1	0.45	1.27 1.72	1.88	1.58	4.63	3.01	2.47 2.47
137984	1 1	0.66		2.25	1.34	6.05	1.01	
76 <b>7346</b> 149 <b>29</b> 67		1.82	1.80 2.19	1.32 1.49	6.76 6.63	1.74 1.16	0.73	2.47 2.47
342721		6.06	3.30	1.93	5.09	0.69	1.30	2.46
7600:		0.96	4.18	3.61	1.37	1.59	1.56	2.46
823850			2.69	1.51	1.29	5.06	1.76	2.46
74490:			3.34	2.72	1.92	3.17	1.16	2.46
39953			1.68	2.54	1.91	3.15	3.01	2.46
95046			1.50	1.01	7.67	1.11	0.97	2.45
48634			1.85	3.58	1.43	3.51	1.87	2.45
86686			4.47	3.47	1.46	1.44	1.40	2.45
88454			2.76	1.70	1.46	3.06	3.23	2.44
00.754	-1	1 2,224	2., 3	3		2.00		,

Table 1-1

897448	31.17	12.76	3.09	2.67	1.77	3.01	1.67	2.44
950989	2.48	1.02	4.36	3.15	1.27	1.86	1.55	2.44
1031583	38.01	15.59	0.37	1.52	1.75	6.11	2.45	2.44
381036	3.54	1.46	2.61	1.83	5.63	1.30	0.81	2.43
109316	12.35	5.08	1.34	0.40	2.84	0.34	7.26	2.43
448117	3.91	1.61	3.94	1.97	3.08	1.50	1.68	2.43
156045	2.01	0.83	3.16	3.14	1.81	2.33	1.71	2.43
884333	4.10	1.69	2.33	3.28	3.72	1.30	1.51	2.43
306771	3.61	1.49	3.02	3.83	1.70	1.84	1.73	2.42
430465	5.55	2.30	0.96	0.93	1.81	7.45	0.92	2.42
123755	5.06	2.10	3.75	1.13	1.03	1.10	5.05	2.41
42803	2.96	1.23	0.60	2.20	1.73	5.20	2.32	2.41
460164	0.80	0.33	2.54	3.13	2.35	3.08	0.95	2.41
1574594	3.08	1.28	4.44	3.10	1.44	1.96	1.07	2.40
141361	1.28	0.54	1.80	0.83	1.31	6.11	1.94	2.40
855336	5.69	2.37	1.65	3.16	2.36	3.39	1.43	2.40
868396	6.77	2.83	5.03	3.73	0.87	1.05	1.28	2.39
627053	11.04	4.64	1.29	2.07	5.16	2.01	1.37	2.38
815772	1.44	0.61	3.35	1.86	3.10	1.62	1.96	2.38
897547	15.69	6.62	3.93	3.84	2.46	0.78	0.84	2.37
202154	29.46	12.46	0.97	2.61	0.99	5.43	1.83	2.37
511107	7.14	3.02	0.73	1.70	3.61	3.54	2.25	2.36
810408	9.85	4.18	1.91	1.77	1.69	3.08	3.34	2.36
757222	5.14	2.18	2.47	6.61	1.12	0.68	0.91	2.36
810734		1.05	3.06	1.80	1.50	4.04	1.32	2.35
1493390		1.53	3.30	1.07	2.47	3.35	1.54	2.34
284865	2.87	1.23	3.64	3.21	1.66	1.49	1.70	2.34
488033	1 1	1.34	2.41	3.80	1.46	3.21	0.82	2.34
37665	• .	0.43	0.55	3.88	2.77	3.16	1.33	2.33
138861		0.74	1.34	3.12	1.63	3.22	2.35	2.33
272049		2.95	3.26	1.35	1.69	4.65	0.71	2.33
427812		4.27	2.58	1.48	5.45	1.15	0.97	2.33
809995		1.21	3.69	1.55	2.04	3.10	1.24	2.32
449275	· '	1.64	5.51	1.68	1.24	2.01	1.16	2.32
731337		0.32	3.30	3.09	1.35	1.91	1.95	2.32
854570		1.78	4.79	3.45	0.94	1.54	0.83	2.31
767988		1.58	1.57	3.20	3.21	1.89	1.63	2.30
455271		0.72	5.02	2.60	1.17	1.48	1.22	2.30
130136		0.13	1.17	1.08	0.84	0.80	7.57	2.29
757191		2.66	1.09	0.88	5.46	2.03	1.99	2.29
266500			1.54	3.23	1.95	3.65	1.08	2.29
72550			3.35	2.31	1.33	3.39	1.03	2.28
155552			1.80	6.52	1.30	0.93	0.86	2.28
62949	4		4.45	3.42	1.09	1.19	1.23	2.28
78140			1.40	1.34	2.90	0.55	5.16	2.27
63213			4.14	1.33	3.68	1.07	1.12	2.27
29383			1.87	1.37	5.66	1.25	1.18	2.26
5056	L		6.49	1.87	0.28	0.43	2.21	2.26
24378			3.68	3.50	1.79	1.35	0.94	2.25
20916			5.12	1.17	2.88	0.92	1.17	2.25
25172				3.39	1.73	3.26	1.65	2.25

Table 1-1

204546	1 10	0.40	1.00	2.07	0.05	4.00	. o.l	224
284545	1.10	0.49	1.06	3.27	0.97	4.66	1.26	2.24
504290	50.88	22.72	1.27	1.74	5.93	1.07	1.18	2.24
71672	4.77	2.14	3.51	1.66	3.03	1.97	0.99	2.23
279394	3.18	1.42	1.37	1.39	5.66	0.95	1.78	2.23
309864	2.49	1.12	8.06	0.69	0.90	0.79	0.67	2.22
291394	6.46	2.92	1.70	1.25	5.58	1.18	1.37	2.22
289645	0.92	0.42	1.62	1.02	1.30	5.62	1.50	2.21
460218	2.10	0.95	1.71	1.76	3.37	3.35	0.86	2.21
856174	0.25	0.11	3.70	4.65	0.88	0.91	0.92	2.21
767784	8.86	4.05	5.50	1.03	1.73	1.72	0.97	2.19
826459	12.36	5.65	0.94	3.16	1.94	3.18	1.72	2.19
141336	2.12	0.97	0.78	0.91	7.64	0.64	0.94	2.18
122239	6.50	2.98	4.86	3.17	0.88	0.92	1.06	2.18
840698	7.45	3.42	3.42	3.27	1.39	1.58	1.22	2.18
303139	2.17	1.00	0.98	0.85	2.90	5.31	0.82	2.17
277761	2.43	1.12	3.26	1.81	1.46	3.22	1.08	2.17
471641	4.79	2.22	0.68	4.45	0.89	4.15	0.64	2.16
470114	3.09	1.44	2.04	3.03	1.58	3.16	0.97	2.15
33045	0.63	0.29	5.38	1.08	1.25	1.95	1.09	2.15
84078	0.34	0.16	0.79	0.75	3.03	4.97	1.17	2.14
111389	2.59	1.21	3.22	1.26	1.20	3.23	1.78	2.14
1536991	10.79	5.05	3.61	1.25	3.84	1.06	0.92	2.14
770935	86.21	40.44	0.25	2.50	1.55	3.29	3.08	2.13
281191	7.36	3.45	3.30	1.86	3.23	0.81	1.46	2.13
149895	6.05	- 2.84	1.33	1.44	6.25	0.94	0.69	2.13
648046	15.56	7.32	0.56	0.94	3.27	2.53	3.34	2.13
855236	2.75	1.30	1.11	1.65	1.52	5.25	1.06	2.12
<b>7</b> 40780	4.71	2.25	1.78	6.08	0.90	0.38	1.35	2.10
844768	6.47	3.08	4.79	3.24	0.75	0.87	0.85	2.10
811097	2.59	1.23	5.78	1.43	0.76	1.89	0.62	2.10
731255	2.04	0.98	1.12	1.83	1.35	5.14	1.01	2.09
950607	19.79	9.50	2.04	1.14	5.08	1.05	1.11	2.08
1472150	35.19	16.93	3.00	1.41	1.13	3.73	1.12	2.08
52338	2.91	1.41	3.16	1.49	4.05	0.57	1.07	2.07
781061	0.73	0.35	0.76	1.88	3.32	3.43	0.88	2.05
1613940		1.70	3.02	3.14	0.88	2.54	0.68	2.05
1604717	13.38	6.59	1.03	1.41	5.95	0.92	0.85	2.03
151477		1.21	0.96	1.86	0.96	5.73	0.57	2.02
563444	6.23	3.10	1.64	3.21	0.69	3.64	0.86	2.01
827168		3.24	1.91	0.82	5.31	1.07	0.95	2.01
845780		2.97	4.42	3.23	0.81	0.67	0.89	2.00
743574			4.33	3.22	0.89	0.69	0.88	2.00
431597			1.67	0.89	5.75	0.72	0.97	2.00
740941			6.11	1.20	0.62	0.67	1.29	1.98
343688			1.39	1.27	5.47	0.79	0.88	1.96
75059			3.00	1.01	1.04	3.21	1.50	1.95
782513			1.16	0.84	5.53	1.41	0.75	1.94
36681			5.26	1.35	1.25	0.46	1.38	1.94
13291			3 <b>.07</b>	1.21	3.40	0.74	1.22	1.93
28289			1.03	1.04	5.87	0.90	0.81	1.93
51211	6 4.59	2.41	1.08	0.94	5.62	1.08	0.80	1.90

291706	2.41	1.27	1.34	1.06	0.98	0.91	5.20	1.90
50114	0.83	0.44	0.97	1.06	0.39	3.57	3.45	1.89
450464	3.97	2.13	3.23	1.41	3.18	0.56	0.95	1.86
626462	5.86	3.24	3.91	3.03	0.70	0.59	0.81	1.81
867717	5.61	3.15	3.27	3.01	0.87	0.68	1.08	1.78
462953	3.21	1.90	0.67	5.53	1.05	0.63	0.57	1.69
41287	0.72	0.45	0.73	0.95	0.53	5.14	0.61	1.59
122428	2.03	1.29	5.48	0.55	0.63	0.63	0.58	1.57

Table 1-2

_					able 1-2			
Clone	Aver of	Aver. of	MPM-115 /	MPM-116/	MPM-117/	MPM-118/	MPM-119/	Aver.
	Tumors	Normals	Aver	Aver	Aver		Aver Normal	Fold
788180	25.18	0.81	62.98	3.35	9.48	55.67	24.72	31.24
646037	24.42	1.02	38.80	10.71	20.96	13.10	35.99	23.91
1034473	33.61	1.51	39.30	2.42	16.40	18.87	34.35	22.27
281003	128.77	6.30	25.82	30.44	5.57	24.33	15.96	20.43
7734 <b>7</b> 9	12.32	0.65	24.85	24.44	9.41	16.91	18.67	18.85
823940	22.39	1.32	15.54	10.31	45.60	5.40	8.09	16.99
281039	14.05	0.94	1.73	6.90	1.46	63.60	1.27	14.99
46166	13.13	0.93	2.81	31.75	16.71	15.27	4.05	14.12
42415	6.86	0.51	4.67	6.71	15.78	1.60	38.87	13.52
291255	15.20	1.35	10.61	10.39	17.35	9.54	8.56	11.29
308682	8.99	0.87	8.35	12.44	6.96	9.39	14.72	10.37
450049	18.37	1.83	8.45	6.05	7.47	5.21	23.07	10.05
283312	63.94	6.63	11.46	5.48	5.23	11.97	14.09	9.65
432194	10.34	1.11	5.77	11.20	10.61	6.96	12.20	9.35
35626	9.42	1.09	6.10	9.04	16.67	3.16	8.40	8.67
222157	8.80	1.02	13.68	8.34	7.21	3.07	11.05	8.67
814053	14.66	1.70	8.20	1.30	6.73	19.59	7.22	8.61
811867	24.57	2.97	9.33	7.93	6.31	11.68	6.18	8.28
35058	15.59	1.89	3.32	10.65	5.05	16.86	5.29	8.24
971279	54.34	6.63	8.84	8.00	7.58	8.74	7.81	8.20
293635	26.47	3.44	5.13	10.21	7.80	7.55	7.78	7.69
32160	26.80	3.59	6.47	7.13	8.59	6.37	8.77	7.47
754280	23.83	3.20	15.06	7.80	8.20	3.02	3.20	7.46
282089	11.18	1.51	1.86	11.16	1.40	21.82	0.82	7.41
745347	2.15	0.29	3.74	8.94	8.54	8.80	6.91	7.39
67401	1.47	0.20	1.25	1.47	30.35	2.21	1.11	7.28
293651	13.35	1.84	3.60	7.75	11.33	6.53	7.12	7.26
784168	B .	7.37	2.50	6.64	8.80	8.22	9.84	7.20
811603		2.74	8.80	6.79	5.12	7.74	7.36	7.16
486035	1	3.47	5.79	11.37	4.91	5.88	7.72	7.13
809603	1	6.94	5.63	5.54	6.31	7.30	10.54	7.06
1048810		3.04	7.11	10.78	4.14	7.28	3.86	6.63
756600	•	7.13	6.98	3.61	3.71	8.73	9.97	6.60
772304		18.10	7.98	4.62	4.60	5.98	8.08	6.25
1631713		0.93	5.38	6.95	7.45	5.29	5.61	6.13
838366			4.41	5.92	6.20	5.88	8.19	
757435			3.80	3.09	17.09	3.11	3.46	
1556526		1	0.88	2.02	1.15	24.74	1.00	
450307				8.38	5.22	1.77	2.92	
796161							5.40	5
1034738		1	t .					
49302								1
292515	1							
810220			1					
207029								
415700 449112								•
53102								1
3627 <b>5</b> :								
50988°								1
19190								
87783		1	•					
01103.	21 23.00	· 1 -4.54	7.48	3.06	4.32	5.71	6.02	5.32

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				Tab	le 1-2			
771317	9.44	1.78	4.49	3.90	5.78	4.69	7.70	5.31
549101	12.81	2.43	7.81	3.40	3.06	5.03	7.02	5.26
84955	5.26	1.01	8.05	6.43	6.35	2.52	2.71	5.21
461670	8.90	1.72	3.21	6.57	7.68	3.75	4.73	5.19
197525	9.52	1.85	2.67	8.08	4.07	6.57	4.40	5.16
730739	36.49	7.10	4.49	4.01	10.50	3.62	3.06	5.14
240694	4.53	0.90	4.65	7.82	3.10	3.41	6.08	5.01
593185	28.06	5.61	1.33	3.36	3.94	10.92	5.45	5.00
212165	41.96	8.42	7.66	4.81	6.54	2.71	3.22	4.99
268211	3.43	0.70	4.14	3.18	3.13	2.23	11.97	4.93
233419	5.55	1.14	3.13	6.59	6.63	2.94	5.09	4.87
296788	3.27	0.67	4.84	4.90	7.08	2.63	4.79	4.85
810558	14.49	3.00	3.18	4.47	4.01	4.93	7.57	4.83
264895	7.86	1.63	3. <b>0</b> 7	6.53	4.76	3.25	6.52	4.82
51773	11.48	2.40	5.99	3.55	3.22	7.62	3.50	4.77
191603	22.14	4.64	6.96	6.89	2.33	3.85	3.81	4.77
417908	12.10	2.57	9.22	5.41	4.37	1.99	2.57	4.71
50188	10.75	2.29	5.42	4.91	4.42	3.76	4.92	4.69
284701	33.05	7.05	3.72	4.93	2.53	9.40	2.86	4.68
416436	11.08	2.37	7.73	5.37	3.95	4.54	1.77	4.67
321470	7.74	1.68	4.78	6.01	4.56	2.65	5.05	4.61
449275	3.81	0.83	10.88	3.32	2.44	3.97	2.28	4.58
263697	4.23	0.93	5.22	5.51	7.03	1.91	3.18	4.57
266106	20.68	4.53	4.04	4.70	9.22	2.05	2.82	4.57
194364	5.44	1.19	2.81	3.35	9.83	4.15	2.64	4.56
259374 877641	10.51	2.33	4.05	2.96	1.50	8.52	5.48	4.50
283173	67.93 17.85	15.10 3.97	8.40	2.61	1.99	5.49	3.99	4.50
868282	8.99	2.01	10.94	2.48	2.73	4.55	1.75	4.49
770066	11.43	2.60	4.17 1.11	4.41 1.71	5.09 4.03	4.45	4.28	4.48
950688	52.65	11.99	5.18	2.58	3.06	13.55 4.94	1.61 6.20	4.40 4.39
490178	24.55	5.63	2.55	2.07	13.03	1.58	2.57	4.36
30428	4.29	0.99	0.90	2.03	2.95	11.84	4.01	4.35
210405	34.02	7.84	3.39	3.97	5.43	4.21	4.69	4.34
795907	5.39	1.25	1.95	2.24	4.60	0.83	11.91	4.31
47542		1.88	6.90	3.68	2.63	4.23	4.04	4.30
297439	48.60	11.33	4.18	7.63	3.83	1.95	3.86	4.29
1472689	5.15	1.21	3.31	0.99	2.88	11.98	2.13	4.26
897767	2.07	0.49	3.67	3.81	4.12	3.26	6.41	4.25
289891	7.82	1.84	4.50	2.99	9.38	2.41	1.95	4.25
853968	12.10	2.86	5.03	5.67	7.48	1.76	1.23	4.23
840788		5.46	4.18	3.11	1.97	7.77	4.07	4.22
856447		2.09	9.08	2.15	2.21	4.88	2.79	4.22
346889	1	0.21	2.54	5.10	1.70	6.26	5.45	4.21
321706		1.86	1	0.98	9.88	1.19	3.34	4.20
855749	Li .	12.03	5.65	3.70	3.56	3.56	4.54	4.20
1031583		9.06		2.62	3.01	10.51	4.21	4.20
785913				3.95	7.94	2.07	2.99	4.19
746152			1	4.05	7.60	1.47	0.48	4.18
845363	1		i	3.53	2.38	5.97	4.42	4.18
377246	•		3	2.77	11.38	1.14	1.90	4.17
129024				0.75	2.32	2.43	7.90	4.13
15026: 82520:		1		5.14	4.58	3.46	3.56	
07770	7 11.96	2.92	3.12	3.72	4.18	5.49	3.94	4.09

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				Tab	le 1-2			
1030855	14.86	3.64	1.14	4.82	2.58	7.86	4.02	4.08
878127	3.53	0.87	2.76	4.48	6.05	4.44	2.62	4.07
138444	4.04	1.00	3.68	7.21	2.94	4.04	2.44	4.06
768395	14.29	3.54	5.36	4.80	4.35	3.14	2.53	4.03
788541	12.11	3.01	2.87	3.90	7.56	1.94	3.89	4.03
1468260	4.53	1.13	7.56	0.78	4.31	2.33	5.15	4.03
787938	5.09	1.27	1.46	4.56	6.98	5.35	1.72	4.01
258300	18.15	4.53	2.52	4.17	7.79	0.73	4.84	4.01
208082	7.90	1.98	3.29	6.85	3.52	3.30	3.00	3.99
785910	4.01	1.01	3.80	2.26	9.32	1.33	3.21	3.98
287728	2.76	0.70	0.75	5.13	1.25	11.82	0.89	3.97
687297	12.18	3.12	4.57	4.32	4.27	2.73	3.63	3.90
741067	64.30	16.50	0.91	4.38	2.37	7.31	4.51	3.90
1031598	5.18	1.34	0.69	2.70	2.63	9.79	3.58	3.88
366815	71.61	18.49	10.51	2.69	2.49	0.91	2.76	3.87
323474	1.25	0.32	6.77	4.12	2.60	2.99	2.88	3.87
288695	3.41	0.88	4.88	4.64	5.25	3.06	1.48	3.86
204111	3.38	0.87	3.40	4.11	3.20	3.52	5.07	3.86
242952	16.94	4.39	4.50	3.65	3.98	4.16	2.99	3.86
1631747	5.09	1.33	5.74	<b>6.7</b> 6	2.76	2.44	1.49	3.84
48286	2.34	0.61	3. <b>34</b>	2.74	4.95	3.47	4.67	3.83
745503	42.62	11.14	2.97	4.02	3.39	3.08	5.68	3.83
744846	2.69	0.70	4.56	. 5.57	3.31	4.45	1.24	3.83
454771	2.69	0.70	7.92	2.22	2.30	1.94	4.70	3.82
233464	14.24	3.73	2.79	2.89	5.83	3.29	4.26	3.81
250869	123.92	32.54	10.99	2.78	2.27	1.03	1.97	3.81
842925	4.61	1.21	3.52	3.54	4.61	4.62	2.67	3.79
843094	49.18	13.00	4.51	3.53	3.71	3.00	4.17	3.78
587415	12.98	3.44	6.76	3.20	6.42	1.08	1.39	3.77
33715	1.36	0.37	2.84	3.84	4.39	3.91	3.66	3.73
75044	3.46	0.93	4.59	3.85	2.61	3.71	3.84	3.72
1467207 366132	3.68	0.99	2.51	4.57	4.41	2.93	4.17	3.72
813973	6.22 3.24	1.68	4.94	2.88	2.87	4.12	3.77	3.72
814017		0.88 23.42	3.24 5.13	3.78 5.73	4.03	3.23 1.59	4.23	3.70 3.68
785616	1	3.12	4.63	3.13 4.47	3.05 3.42	3.47	2.90 2.37	3.67
42803		0.81	0.91	3.34	2.63	7.89	3.52	· 3.66
488964	1	2.13	11.81	2.73	1.04	1.76	0.88	3.64
151055		7.51	3.65	6.52	1.80	2.91	3.31	3.64
784258		2.77	4.79	2.92	5.87	1.93	2.67	3.64
81331		0.78	1.10	2.95	0.88	12.05	1.18	3.63
897042		1.36	6.51	3.85	2.87	3.80	1.14	3.63
771048		2.90	4.89	3.20	3.32	2.85	3.91	3.63
795265		1.26	1.26	2.63	2.84	9.65	1.76	3.63
753252		2.14	2.88	3.17	3.38	4.62	4.10	3.63
430465		1.53	1.44	1.39	2.71	11.18	1.38	3.62
429678	•		3.79	2.41	5.14	4.64	2.08	3.61
47384		1.17	2.95	3.72	4.97	3.42	2.98	3.61
34227			3.73	1.46	4.16	6.29	2.35	3.60
712950			0.83	1.53	3.05	11.90	0.65	3.59
24554	1		1.33	6.79	3.91	4.02	1.91	3.59
147598			4.18	2.40	3.01	6.04	2.26	3.58
35885			6.12	3.37	2.69	1.80	3.82	3.56
4532	7 8.18	2.30	2.35	3.39	3.66	4.62	3.78	3.56

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				Te	ble 1-2			
950410	3.18	0.89	1.81	7.55	3.00	2.68	2.76	3.56
1592006	5.30	1.51	4.10	3.01	3.34	3.68	3.47	3.52
741474	4.93	1.40	2.90	2.23	3.62	5.57	3.24	3.51
292042	9.39	2.67	3.33	4.22	4.13	3.14	2.73	3.51
280217	20.52	5.86	4.76	0.46	1.34	10.23	0.72	3.50
42636	1.11	0.32	2.78	3.16	4.86	2.57	4.13	3.50
550141	78.20	22.40	5.62	2.32	2.57	3.16	3.79	3.49
159608	14.43	4.14	6.82	1.07	1.06	5.09	3.39	3.49
383185	12.04	3.46	6.25	2.54	6.69	0.98	0.94	3.48
491184	6.35	1.83	4.07	1.43	10.01	0.85	1.00	3.47
786525	1.46	0.42	2.15	1.29	2.53	10.04	1.31	3.46
305538	11.44	3.31	4.78	4.20	1.98	1.85	4.49	3.46
415437	3.52	1.02	2.69	3.45	3.10	4.16	3.88	3.46
259350	22.84	6.64	2.91	2.83	2.97	3.77	4.71	3.44
950709	4.17	1.22	5.93	2.37	2.11	4.13	2.61	3.43
141505	6.38	1.86	2.88	4.20	3.06	2.08	4.90	3.42
109200	3.17	0.93	5.89	2.59	3.19	2.96	2.46	3.42
1390584	6.54	1.91	1.98	3.12	3.26	8.11	0.63	3.42
823819	11.96	3.51	4.50	4.09	2.11	2.94	3.43	3.41
272750	20.57	6.04	3 <b>.75</b>	5.31	2.77	2.65	2.54	3.40
246869	4.68	1.38	4.93	2.40	3.61	2.07	3.98	3.40
784253	11.76	3.47	2.81	4.36	3.28	2.90	3.61	3.39
490766	3.48	1.03	4.47	3.60	2.64	2.91	3.27	3.38
356863	3.18	0.94	4.17	3.83	2.02	3.24	3.63	3.38
837904	103.13	30.63	2.21	2.23	2.11	4.52	5.77	3.37
462468	3.24	0.96	3.67	4.25	3.86	1.95	3.06	3.36
121997	5.10	1.52	4.66	2.43	5.54	1.60	2.55	3.36
700900	9.94	3.00	3.26	2.88	3.29	3.75	3.40	3.31
26616	1.94	0.59	4.02	2.79	3.28	3.05	3.40	3.31
1589973	5.17	1.57	2.96	3.72	2.30	5.22	2.31	3.30
392678	34.95	10.61	2.24	2.72	3.38	3.80	4.32	3.29
586725	2.32	0.70	3.69	2.64	2.18	3.31	4.64	3. <b>29</b>
121489	34.70	10.55	3.98	4.08	3.24	2.67	2.47	3.29
503851	18.15	5.54	1.87	3.27	5.43	2.90	2.90	3.28
854874	2.22	0.68	2.50	3.02	5.99	2.47	2.32	3.26
877832 253009	44.43 115.86	13.65	3.45	3.31	2.92	3.09	3.50	3.25
824723		35.75	1.83	3.37	1.51	5.38	4.13	3.24
755409	14.11 7.81	4.37	2.37	2.55	3.40	5.30	2.52	3.23
506564	12.93	2.43 4.03	3.30	3.12	4.05	3.18	2.47	3.22
109153	5.46	1.70	4.44	2.86	3.21	1.98	3.55	3.21
213682		4.10	4.21 0.63	2.85	3.08	2.33	3.55	3.20
133717		2.97	2.19	2.78 2. <del>2</del> 2	2.14 3.36	7.22	3.16	3.19
361659		4.23	1.54	3.83	4.60	2.66	5.49	3.18
232628	1	1.27	4.95	2.33	5.56	3.93 1.30	1.99	3.18 3.16
283034		3.05	3.07	5.05	2.86		1.63	
490232		0.85	3.64	3.48	3.06	2.31 1.82	2.46 3.75	3.15
781366	i	3.84	3.48	2.65	1.25	2.22	6.10	3.15 3.14
66317		1.76	6.84	2.02	1.77	4.59	0.10	3.14
359781		2.50	1.84	5.41	2.09	3.84	2.45	3.13
564801		3.96	2.96	1.38	8.87	1.73	0.67	3.13
587595		179.15	0.74	2.46	3.02	6.40	2.98	3.12
39885		0.30	0.92	2.92	4.86	4.97	1.87	3.11
45578			4.85	2.27	2.38	3.60	2.39	3.10
	•	•	•			5.00	5,	

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				Tab	le 1-2			
743749	5.10	1.65	4.98	3.05	2.98	3.05	1.42	3.10
839081	16.60	5.37	5. <b>25</b>	5.49	0.97	2.62	1.11	3.09
109316	12.35	4.01	1.69	0.50	3.59	0.43	9.18	3.08
358083	2.45	0.80	3.44	2.58	4.41	1.95	2.99	3.07
130773	3.71	1.21	2.86	3.98	3.19	2.42	2.91	3.07
610011	34.63	11.27	3.99	2.71	2.59	2.28	3.79	3.07
429574	3.77	1.23	2.74	4.01	3.30	1.01	4.26	3.06
167205	7.85	2.56	2.90	5.74	3.48	1.88	1.31	3.06
825718	4.03	1.32	2.59	2.65	4.00	3.21	2.84	3.06
42659	10.21	3.36	3.63	3.18	3.06	2.09	3.23	3.04
1631682	1.61	0.53	4.08	2.09	3.06	2.23	3.71	3.03
843234	7.02	2.32	3.12	3.14	2.40	2.69	3.80	3.03
290642	7.76	2.56	4.18	2.71	2.32	3.00	2.91	3.02
51700	3.66	1.21	3.02	2.75	3.37	2.08	3.87	3.02
214006	4.67	1.55 4.59	6.05	4.62	1.41	2.11	0.89	3.02
43 1286 78 1454	13.81 9.75	3.24	2.40 2.57	3.23 1.19	3.52 9.17	2.10 1.13	3.80 0.98	3.01 3.01
433253	7.29	2.43	2.31	2.35	2.05	5.26	3.03	3.00
359661	26.24	8.79	2.02	4.95	1.82	3.95	2.19	2.99
811953	17.84	5.98	3.25	3.61	3.69	1.97	2.41	2.98
196345	122.93	41.22	1.48	2.47	2.71	3.86	4.39	2.98
950473	9.36	3.14	2.36	3.35	2.97	2.65	3.56	2.98
46294	1.03	0.35	1.45	2.41	1.44	5.75	3.83	2.97
37980	0.84	0.29	1.32	2.54	1.61	7.51	1.81	2.96
512116	4.59	1.55	1.67	1.46	8.72	1.68	1.24	2.96
137638	5.06	1.71	3.63	2.92	2.17	2.49	3.54	2.95
810753	4.95	1.68	3. <b>05</b>	2.91	3.48	2.96	2.35	2.95
592403	0.46	0.16	0.70	2.31	2.27	6.90	2.56	2.95
273546	9.10	3.09	4.02	3.24	2.32	2.44	2.71	2.95
238 <b>821</b>	6.54	2.22	3.78	1.07	3.22	0.62	6.02	2.94
111812	5.56	1.89	3.45	4.26	4.44	1.20	1.35	2.94
840 <b>894</b>	34.81	11.87	1.12	3.23	2.53	4.70	3.09	2.93
50114	0.83	0.28	1.50	1.65	0.60	5.53	5.35	2.92
971367	35.22	12.06	3.23	1.99	1.70	5.64	2.05	2.92
1473300	12.44	4.26	1.78	2.25	2.22	3.03	5.31	2.92
824704 8976 <b>70</b>		0.70 2.19	3. <b>0</b> 7 3. <del>64</del>	2.49 3.98	3.23 2.74	2.63 2.73	3.15 1.43	2.91 2.90
430153	1	1.54	3.45	3.54	2.88	1.69	2.95	2.90
31225		1.64	2.16	2.29	4.85	2.20	3.01	2.90
743828		1.15	1.18	1.72	5.33	1.00	5.28	2.90
1521977		0.72	3.62	1.94	2.94	3.07	2.92	2.90
511632		2.71	2.19	2.68	1.95	4.09	3.56	2.89
36 <b>0885</b>		13.93	4.07	2.17	4.57	1.22	2.42	2.89
1492238		4.83	3.53	3.13	2.20	3.54	2.04	2.89
784830	1.83	0.63	4.04	1.72	3.99	1.90	2.77	2.88
76 <b>779</b> 8		1.98	2.68	2.09	2.38	3.81	3.45	2.88
840590		7.54	2.19	1.06	2.15	7.48	1.51	2.88
878316		1.86	2.30	2.73	1.60	4.36	3.36	2.87
725489		9.34	3.59	3.28	2.57	3.29	1.57	2.86
20916		1.15	6.50	1.48	3.65	1.17	1.48	2.86
48843			4.07	2.58	2.02	1.74	3.84	2.85
73130			3.77	3.06	2.31	1.95	3.15	2.85
56 <b>523</b> :			1.31	3.41	1.55	7.11	0.85	2.85
50429	50.88	17.89	1.61	2.21	7.54	1.36	1.50	2.84

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	Table 1-2									
823588	7.72	2.72	2.42	3.62	3.20	2.49	2.44	2.83		
878544	19.20	6.78	5.95	1.60	3.38	1.81	1.41	2.83		
26736	9.52	3.36	3.45	0.77	0.57	6.03	3.33	2.83		
145388	2.66	0.94	3.12	3.26	4.21	2.28	1.26	2.83		
76196	2.53	0.90	1.68	2.14	2.06	3.22	5.03	2.83		
624271	5.74	2.03	3.08	2.14	1.61	6.02	1.25	2.82		
66902	3.21	1.14	3.84	1.85	1.41	0.79	6.22	2.82		
86160	2.10	0.74	2.49	3.24	2.74	2.51	3.12	2.82		
526945	1.14	0.40	3. <b>22</b>	1.77	2.51	3.16	3.41	2.82		
131621	10.21	3.63	3.64	1.77	4.66	1.90	2.11	2.82		
815036	1.62	0.58	3.61	3.75	2.37	3.01	1.33	2.81		
416833	6.07	2.16	1.37	3.86	1.76	4.96	2.11	2.81		
810221	3.85	1.37	1.41	2.35	3.96	2.88	3.43	2.81		
1536991	10.79	3.85	4.74	1.63	5.04	1.39	1.21	2.80		
1240220	7.60	2.71	0.70	2.40	5.33	4.42	1.16	2.80		
125308	2.61	0.93	4.31	2.08	3.64	2.05	1.93	2.80		
502690	4.09	1.46	2.11	3.13	2.39	4.17	2.19	2.80		
1635874	6.06	2.17	5.71	2.90	1.66	2.16	1.54	2.79		
32076	1.03	0.37	1.94	2.97	2.60	3.25	3.21	2.79		
856535	1.40	0.50	3.08	3.52	2.81	2.44	2.11	2.79		
30793	7.89	2.84	2.22	2.49	3.31	3.04	2.85	2.78		
826102	4.19	1.51	1.94	1.78	7.59	1.08	1.50	2.78		
356992	21.95	7.93	3.72	2.43	2.23	3.22	2.25	2.77		
810571	4.37	1.58	1.62	3.67	1.77	4.25	2.52	2.76		
365098	4.93	1.78	4.88	1.79	3.50	1.53	2.10	2.76		
788655	7.66	2.78	2.38	1.89	4.08	3.62	1.83	2.76		
186918	3.04	1.10	1.49	2.82	2.33	5.76	1.40	2.76		
842784	326.54	118.59	0.68	2.49	1.56	5.61	3.44	2.75		
127396	14.07	5.12	2.63	3.54	0.99	4.76	1.83	2.75		
261472	9.31	3.39	2.65	1.79	2.40	5.00	1.89	2.75		
813179	6.60	2.40	1.14	8.50	0.98	2.38	0.73	2.75		
240430 1435029	3.52	1.28	5.57	1.49	1.68	2.99	1.98	2.74		
124824	1.84 30.85	0.67	2.78	3.83	3.22	2.31	1.55	2.74		
1493390		11.28 1.31	2.73 3.84	1.75	1.56	2.43	5.21	2.73		
251936	7.48	2.74	3.40	1.25	2.88	3.90	1.79	2.73		
343609	1.58	0.58	2.30	3.01 2.63	2.19	2.17	2.89	2.73		
125722	3.28	1.21	4.28	2.03	3.03 2.60	2.56	3.13	2.73		
322051	11.58	4.27	2.00	2.69	3.96	3.74 1.51	0.81 3.38	2.72 2.71		
756556		6.47	1.96	2.33	5.07	2.59	1.60	2.71		
430318	,	4.02	1.74	2.76	3.55	1.94	3.54	2.71		
795555		1.25	3.07	3.53	2.44	2.31	2.14	2.70		
951125		7.00	2.18	5.16	1.53	2.32	2.28	2.69		
843150		1.90	2.80	3.47	2.10	1.72	3.38	2.69		
50772	18.10	6.74	1.30	2.66	2.08	5.85	1.54	2.68		
80500	140.81	52.48	0.85	2.49	1.54	4.92	3.61	2.68		
137189	5.26	1.96	1.82	3.19	2.48	3.67	2.23	2.68		
279482			2.09	1.33	7.28	0.95	1.69	2.67		
745103		3.33	1.82	3.08	3.04	1.84	3.55	2.67		
760148	1	4	2.27	2.03	3.48	3.23	2.28	2.66		
823615	4		3.54	1.46	5.77	1.58	0.93	2.65		
783645	1		3.48	2.53	3.22	1.85	2.18	2.65		
81578	I.	. 1		2.44	4.22	1.59	1.60	2.65		
785793	55.68	21.05	3.09	2.27	3.82	1.77	2.28	2.65		

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				Tabk	: 1-2			
277660	14.16	5.37	1.96	2.26	2.29	3.29	3.39 2.	.64
66336	1.67	0.63	4.14	4.60	4.12	0.16		.63
1630990	20.64	7.84	4.29	2.18	1.71	4.51		.63
247609	3.15	1.20	2.53	1.52	2.16	3.23	3.74 2	.63
898123	4.77	1.81	1.92	3.20	3.04	1.92	3.07 2	.63
40136	5.23	1.99	1.90	1.77	2.01	1.67	5.77 2	.62
46367	1.63	0.62	1.73	1.61	1.24	7.45	1.09 2	.62
281072	5.02	1.92	1.08	2.04	1.92	5.46	2.57 2	62
1606837	23.76	9.09	3.80	1.46	1.83	3.18		.61
308726	14.73	5.64	3.93	2.00	4.67	0.90		2.61
784116	0.26	0.10	2.07	3.32	3.40	2.11		2.61
46620	5.25	2.02	1.48	2.20	1.83	3.69		2.61
504544	12.31	4.73	1.63	2.83	1.76	5.44		2.60
279394	3.18	1.23	1.59	1.61	6.58	1.11		2.59
262865	30.48	11.78	3.45	2.03	4.88	0.98		2.59
123980	2.38	0.92	1.80	3.18	2.76	1.49		2.59
293830	4.43	1.72	2.14	1.56	6.45	1.42		2.58
383188	2.79	1.08	3.03	2.15	1.40	2.92		2.58
853998	12.39	4.81	1.79	2.00	3.76	2.32	1	2.58
813651	4.96	1.93	1.72	2.40	2.48	3.04		2.58
203350	1.91	0.74	2.76	1.43	3.30	3.24		2.57
240961	2.11	0.82	2.13	4.67	3.19	1.71		2.57
1035765		2.41	2.60	0.83	3.16	0.47	5.78	2.57
970271	3.16	1.23	2.05	3.35	2.83	1.20	3.37	2.56
950836	34.79	13.60	2.31	1.27	6.82	0.77	1.62	2.56
461699	3.44	1.35	1.58	2.10	5.96	1.28	1.84	2.55
743851	8.22	3.22	3.05	2.50	2.28	3.39	1.54	2.55
731290	11.60	4.56	1.16	2.19	1.46	7.52	0.38	2.54
1475842	224.84	89.33	0.55	1.42	1.77	6.27	2.56	2.52 2.51
796513		7.09	3.08	2.43	2.08	3.84	1.14 3.25	2.50
8290		0.48	2.88	1.65	1.67	3.07	5.22	2.49
12375	· ·		3.88	1.17	1.06	1.14 0.91	0.70	2.49
88460			2.43	1.59	6.84	2.19	3.33	2.48
32361			3.60	1.64	1.66 1.35	4.43	1.33	2.47
147215			3.57	1.68	3.55	1.12	1.25	2.47
95057			3.24 1.32	3.20 3.06	3.37	2.76	1.82	2.47
34505		1 1	1.32	2.00	5.71	2.00	0.89	2.47
48749			1.73	2.50	5.68	1.17	1.01	2.45
28907 160434		1	1.10	1.51	0.94	6.33	2.34	2.45
89691			3.11	4.47	0.31	3.53	0.78	2.44
25133			1.73	1.78	2.05	3.30	3.33	2.44
47011			2.30	3.43	1.78	3 <b>.57</b>	1.10	2.44
8546°			1.30	7.14	1.43	0.90	1.41	2.44
2458			4.73	1.10	4.77	0.61	0.96	2.43
284			1.13	1.64	1.72	5.29	2.37	2.43
3445			1.02	3.29	3.08	2.11	2.65	2.43
5119		4	•	2.42	0.61	7.65	1.07	2.43
8981	1			2.53	3.20	1.41	3.07	2.42
2917				1.35	1.24	1.16	6.59	2.41
	21			691	0.97	0.98	1.86	2.40

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200599 13.42

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3.50

6.17

5.61

25**8521** 

1412481

1475881

		Table 1-2										
252904	5.40	2.26	3.71	1.41	3.30	1.76	1.77	2.39				
430614	2.38	1.00	2.34	3.41	1.17	4.02	0.99	2.39				
381036	3.54	1.49	2.56	1.79	5.52	1.27	0.79	2.38				
448117	3.91	1.64	3.86	1.93	3.01	1.46	1.64	2.38				
250673	2.66	1.12	5.54	1.03	1.65	1.42	2.21	2.37				
593929	17.74	7.51	3.61	1.43	5.14	0.93	0.70	2.36				
841615	12.16	5.15	2.59	1.62	5.18	1.30	1.11	2.36				
52338	2.91	1.24	3.59	1.69	4.60	0.65	1.22	2.35				
137254	1.35	0.58	3.41	1.79	3.95	1.23	1.38	2.35				
767994	4.29	1.83	2.20	1.22	7.00	0.55	0.73	2.34				
197765	8.83	3.78	2.35	0.96	6.16	0.88	1.33	2.34				
254310	10.26	4.39	2.47	1.99	5.33	1.07	0.80	2.33				
855244	15.78	6.78	4.53	0.61	5.42	0.42	0.66	2.33				
144926	9.20	3.96	3.10	1.65	2.25	3.17	1.43	2.32				
565779	31.62	13.68	3.23	1.26	1.97	3.57	1.53	2.31				
563451 462325	4.17	1.82	3.24	2.08	3.69	1.02	1.43	2.29				
	8.76	3.83	1.92	0.60	1.72	5.27	1.93	2.29				
132911	52.56	22.96	3.65	1.43	4.04	0.88	1.45	2.29				
827168 1070015	6.51 3.26	2.85	2.17	0.93	6.04	1.22	1.08	2.29				
293243	13.17	1.43 5.84	0.69	2.05	0.53	3.30	4.86	2.28				
743211	3.84	1.71	3.23 1.36	1.48 1.66	4.46 5.64	0.69	1.43	2.26				
259462	3.69	1.65	0.66	0.89	0.75	1.28 8.24	1.27	2.24				
291345	70.40	31.69	5.27	1.32	2.52	0.90	0.65	2.24				
281191	7.36	3.32	3.42	1.94	3.36	0.84	1.10	2.22 2.22				
362080	3.81	1.72	1.00	0.91	4.67	3.13	1.35	2.21				
825058	3.42	1.55	2.14	0.93	6.34	1.08	0.54	2.21				
290841	6.35	2.89	3.60	3.33	1.43	1.62	1.02	2.20				
291057	1.34	0.61	1.18	1.53	3.04	4.12	1.04	2.18				
855624	17.75	8.17	0.67	0.78	0.54	0.73	8.15	2.17				
50344	0.85	0.39	0.80	0.85	0.73	7.58	0.85	2.16				
1056172	10.33	4.78	2.40	1.52	0.48	5.66	0.74	2.16				
1056203	1.92	0. <b>89</b>	1.05	0.95	1.43	6.28	1.07	2.15				
78 <b>6609</b>	4.81	2.23	0.54	0.55	1.98	7.08	0.62	2.15				
248261	1.51	0.70	3.67	1.05	0.77	0.89	4.36	2.15				
429047	3.17	1.48	1.32	2.03	3.22	1.07	3.04	2.14				
744933		1.01	0.80	0.91	0.71	7.21	1.06	2.14				
461955	1	0.91	3.10	3.01	1.55	1.32	1.68	2.13				
81484	1	5.15	1.73	3.05	2.09	0.66	3.02	2.11				
897547		7.46	3.48	3.41	2.18	0.69	0.75	2.10				
796366		17.03	3.14	1.59	3.66	0.93	0.94	2.05				
141336 1291658		1.03	0.73	0.85	7.18	0.60	0.88	2.05				
841302	•	1.00	3.72	1.14	3.32	0.94	1.12	2.05				
627688		3.64 3.62	1.25	1.26 1.30	5.92	0.84	0.91	2.04				
811097		1.28		1.38	5.58	1.43 1.83	0.63 0.60	2.03 2.02				
361943		1.43		1.71	0.73 <b>5.64</b>	0.71	0.95	2.02				
75059		10.94		1.03	1.07	3.28	1.53	2.00				
28289		1.96	4	1.07	6.06	0.92	0.83	1.99				
1492967			1	1.20	5.34	0.93	0.83	1.99				
22600				5.49	0.87	1.20	1.00	1.99				
11985	I.			3.58	0.65	0.63	0.64	1.98				
78428				1.06	3.08	1.24	1.07	1.98				
14079				1.05	5.74	0.76	1.05	1.96				
	•	-	-				_	•				

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Table	4 2
3 BURE	1-2

704076 142326 840944 377672	5.70 1.49 79.86 2.59	2.95 0.78 42.51 1.39	1.28 5.18 3.82 0.54	1.04 1.78 0.54 0.57	3.20 0.80 4.16 0.40	3.25 0.91 0.61 7.29	0.88 0.92 0.27 0.56	1.93 1.92 1.88 1.87 1.85
431597 32381 471641 753794	5.45 2.18 4.79 1.65	2.95 1.18 2.74 0.94	1.55 0.60 0.55 0.66	0.82 0.99 3.60 1.32	5.32 3.26 0.72 0.68	0.67 1.10 3.36 5.31	0.90 3.29 0.52 0.78	1.85 1.75 1.75
950464 289832 309864 1492202	9.75 6.67 2.49	5.67 4.11 1.62 1.98	1.05 1.15 5.57 0.42	0.71 0.75 0.48 0.35	5.37 5.01 0.62 0.43	0.77 0.74 0.55 0.34	0.68 0.48 0.47 5.02	1.72 1.62 1.54 1.31

Table 1-3

	Lable 1-3									
I	GenBank	Aver. of	Aver. of	Aver.	Tumor/	Tumor /	T/N	T/N	T/BPH	T/BPH
Clone	Acc No.	Tumors	Normals	of BPH	Normal	врн	5X, ≥1/5	3X, <u>&gt;2</u> /5	5X, ≥1/5	3X, <u>&gt;2</u> /5
	AA453562	25.18	0.81	0.50	31.24	50.69	+	+	+.	+
	AA196979	24.42	1.02	0.75	23.91	32.70	• +	+	+	+
1034473	AA779728	33.61	1.51	1.35	22.27	24.97	+	+	+	+
281003	N50880	128.77	6.30	4.90	20.43	26.26	+	+	+	. +
308682	W25202	8.99	0.87	0.27	10.37	32.91	+	+	+	+
281039	N47717	14.05	0.94	0.90	14.99	15.57	+	+	+	+
284701	N64840	33.05	7.05	1.52	4.68	21.68	+	+	+	+
823940	AA490213	22.39	1.32	2.83	16.99	7.91	+	+	+	+
42415	R60981	6.86	0.51	0.72	13.52	9.49	+	+	+	+
46166	H09076	13.13	0.93	1.71	14.12	7.68	+	+	+	+
	AA432103	53.04	7.37	3.83	7.20	13.84	+	+	+	+
	AA427899	12.32	0.65	5.99	18.85	2.05	+	+	_	
222157		8.80	1.02	0.72	8.67	12.14	+	+	+	+
	AA043109	24.75	3.47	1.86	7.13	13.32	+	+	+	+
	AA703396	18.37	1.83	1.80	10.05	10.21	+	+	+	· +
66336		1.67	0.63	0.10	2.63	16.70		+	+	+
35058		15.59	1.89	1.44	8.24	10.81	+	+	+	+
26736		9.52	3.36	0.62	2.83	15.37	+	+	+	+
	AA159714	28.06	5.61	2.29	5.00	12.23	+	+	+	+
32160		26.80	3.59	2.77	7.47	9.69	+	+	+	+
	AA465495	14.66	1.70	1.75	8.61	8.38	\	+	+	+
292515		16.22	2.86	1.75	5.67	10.38	ļ <del>,</del>	+	+	+
	AA436097	36.49	7.10	3.45	5.14	10.57	+	+	+	+
282089		11.18	1.51	1.41	7.41	7.92	+	+	+	+
283312	1	63.94	6.63	12.14	9.65	5.27	+	+	+	+
291255		15.20	1.35	4.65	11.29	3.27	+	+	+	+
	AA621342	20.15	3.04	2.59	6.63	7.7 <b>8</b>	1	+	+	+
197525	1	9.52	1.85	1.06	5.16	9.01	+ +	+	+	+
67401	L		0.20	0.23	7.28	6.27	+	-	+	•
	AA121266		5.63	2.73	4.36	8.99	1	-	+	- +
	AA935560		0.15	0.12	5.96	7.33			+	•
	AA419486		2.23		4.18	8.70		+	+	+
	AA780190		3.90		5.74	6.82		+	+	+
259374			2.33		4.50	8.04		+	+	+
	AA437224				6.11	6.27		, +	+	. +
	AA479284				7.46	4.85	•	+	+	+
	5 AA625634				5.32	6.90	•	+	+	+
	0 AA282272				3.59	8.58	· ·	+	+	+
5077					2.68	9.47			+	+
	8 A1000271				1.50			_	+	+
	4 AA679414							+		+
	4 AA047260							•	+	+
	1 AA845156							•	+	· <u>-</u>
28317								+	+	+
	7 AA454963							+	_	+
2651	L							•	+	+
	7 AA23522							+	+	+
	4 AA23224							•	+	+
	0 AA88492							+	+	+
25830								+	+	+
	9 AA68290							+	- T	•
	18 AA99185							т	+	+
101044	1014V33193	. 133.3	J J 1.01	5 16.44	2.38	8.2.	2 I -	-	*	τ

				Ta	able 1-3					
461522	AA705237	1.21	0.65	0.14	1.87	8.74	•	<del>-</del> .	+	+
	AA633957				4.48	6.12	`+	+	+	+
280217	N47941	20.52			3.50	7.09	+	+	+	+
240694	H78135	4.53			5.01	5.47	+	+	+	+
321706	W35398	7.80	1.86		4.20	6.23	+	+	+	+
	AA430545	11.43	2.60	1.93	4.40	5.94	+	+	+	+
811603	A1732784	19.62	2.74	6.22	7.16	3.15	+	+	-	+
51773	H23211	11.48	2.40	2.09	4.77	5.50	+	+	+	+
66317	T66815	5.50	1.76	0.78	3.13	7.06	+	+	+	+
35626	•	9.42	1.09	6.51	8.67	1.45	+	+	-	-
	AA682851	9.39	1.62	2.25	5.80	4.17	+	+	+	+
	AA702781	14.83	7.21	1.88	2.06	7.90	-	-	+	+
	AA416843	11.60	4.56	1.57	2.54	7.40	+	-	+	+
	AA504783	4.27	2.51	0.52	1.70	8.20	-	-	+	+
	AA136125	35.83	12.59	5.15	2.85	6.96	+	+	. +	+
	AA481464	47.06	7.13	14.79	6.60	3.18	+	+	•	+
293635	1	26.47	3.44	12.94	7.69	2.05	+	+	-	•
	AA426110	0.86	0.37	0.12	2.32	7.41	-	-	+	+
	5 AA173189	3.58	2.33	0.44	1.54	8.16	-	-	+	+
15105		27.32	7.51	4.62	3.64	5.92	+	+	+	+
29743		48.60	11.33	9.26	4.29	5.25	+	+	+	+
	6 AA457172	11.15	1.82	3.27	6.12	3.41	+	+	-	+
	9 AA677629	0.87	0.46	0.11	1.87	7.58	-	-	+	+
	7 AA173598	3.07	2.45	0.38	1.25	8.17	-	-	+	+
163171	L .	5.68	0.93	1.77	6.13	3.21	+	+	-	+
	1 AA 129551	12.35	3.96	1.99	3.12	6.21	+	-	+	+
29365		13.35	1.84	6.56	7.26	2.04	+	+	-	-
	17 AA625666	2.15	0.29	1.14	7.39	1.88	+	+	•	-
	04 AA404486	113.14	18,10	38.24	6.25	2.96	+	+	•	+
2620:	1	10.75	5.15	1.52	2.09	7.08		-	+	+
813		2.84	0.78	0.51	3.63	5.53	+	-	+	+
	94 AA482340	34.81	11.87	5.66	2.93	6.15	-	+	+	+
	81 AA488732	16.60	5.37	2.78	3.09	5.98	+	+	+	+
4179		12.10	2.57	2.82	4.71	4.29	+	+	+	+
	66 AA459761	34.94	17.03	5.05	2.05	6.92	.   -	+	+	+
	61 AA011095	26.24	8.79	4.39	2.99	5.97	'   -	+	+	+
	188 H17943		2.29	2.53	4.69	4.25		+	-	+
	955 T74846		1.01	1.43	5.21	3.69	)	+	+	+
159			4.14	2.69	3.49	5.36		+	+	+
	584 AA843592		1.91	1.21	3.42	5.41		+	+	+
825	736 AA504842		0.60	0.13	1.54	7.20		-	+	+
	161 T66154		13.40	2.80	1.51	7.2		· -	+	+
	938 AA45227		1.27	1.08	4.01	4.7			+	+
	603 AA44299		6.94	29.52	7.06	1.6			-	-
	109 AA52132		0.54	0.10	1.38	7.2			+	+
	1606 AA63000		13.42	5.43	2.49	6.1			. +	+
	5822 AA49137		9.48		1.48	7.1			+	+
	1028 AA07049		3.89						-	+
	4017 AA45566				3.68			+ +	,	+
	4062 AA46533	18.0 8			1.72			-	+	+
	6103 AA52141	6 1.31			1.61		1	•		+
	1670 AA7799	19 8.90			5.19		;	+ +		+
41	5700 W8569	3.36					1	+ +	· •	+
	5163 AA4811		0.57	7 0.19	2.09	6.	36		•	•
	-									

Page 2

					Table 1-3					
810558	AA464568	14.49	3.00	4.01	4.83	3.61	+	+	+	+
810220	AA464689	2.17	0.39	0.77	5.60	2.82	+	+	-	+
	AA010932	7.82	2.50	1.49	3.13	5.26	+	+	+.	+
	AA451807	4.58	1.26	0.97	3.63	4.74	+	-	+	+
	AA485333	2.49	1.26	0.39	1.97	6.38	-	-	+	+
212165	H68845	41.96	8.42	12.58	4.99	3.34	+	+	+	+
138444	R68272	4.04	1.00	0.95	4.06	4.24	+	+	+	+
	AA488986	14.11	4.37	2.84	3.23	4.96	+	+	+	+
186918	H43317	3.04	1.10	0.56	2.76	5.38	+	-	+	+
191904	H38880	7.67	1.44	2.75	5.32	2.79	+	+	-	+
1630990	A1018613	20.64	7.84	3.79	2.63	5.44	-	+	+	+
	AA608567	52.65	11.99	14.32	4.39	3.68	+	+	+	+
	AA476234	9.44	1.78	3.46	5.31	2.73	+	+ .	-	-
	AA521394	1.20	0.83	0.18	1.44	6.59	-	-	+	+
30428	R42061	4.29	0.99	1.16	4.35	3.68	+	+	+	+
	AA620995	10.33	4.78	1.77	2.16	5.84	+	-	+	+
251806	H96647	2.94	1.83	0.46	1.60	6.39	•	-	+	+
305538	N89861	11.44	3.31	2.54	3.46	4.50	•	+	+	+
52933	H29315	3.59	1.99	0.58	1.80	6.14	•	-	+	+
587415	A1733857	12.98	3.44	3.12	3.77	4.16	+	+	+	+
	AA490522	1.38	1.09	0.21	1.27	6.65	-	-	+	+
416436	t .	11.08	2.37	3.42	4.67	3.24	+	+	+	· +
321470		7.74	1.68	2.36	4.61	3.29	+	+	-	+
	AA486444	4.61	1.21	1.13	3.79	4.07	-	+	-	+
	AA666366	85.06	37.74	15.20	2.25	5.60	-	-	+	+
	AA644092	29.59	7.07	8.08	4.18	3.66	+	+	+	+
	AA018675	16.82	3.11	6.91	5.41	2.43	+	+	-	-
253132	B	1.40	1.21	0.21	1.16	6.66	-	-	+	+
233419		5.55	1.14	1.90	4.87	2.93	+	+	-	+
	AA953229	5.30	1.51	1.24	3.52	4.28	-	+	-	+
	AA608636 AA620591	3.00	1.87	0.48	1.60	6.18	-	-	+	+
358 <b>85</b> 0	4	6.84	6.68	1.01	1.03	6.75	-	-	+	+
	AA427395	21. <b>2</b> 9 10.54	5.98 2.90	5.06 2. <b>58</b>	3.56	4.21	+	+	+	+
	AA486533	79.86	42.51	13.69	3.63 1 <b>.88</b>	4.08 5.83	-	+	++	+
268211		3.43	0.70	1.23	4.93	2.78	- +	+		+
	AA487914	15.85	7.18	2.88	2.21	5.50	-		++	+
	AA461092	30.31	5.27	15.52	5.75	1.95	+	+	<b>T</b>	<del>-</del>
26610		20.68	4.53	6.63	4.57	3.12	+	+	+	+
	2 AA458516	10.13	5.34	1.75	1.90	5.78	'	_	+	+
	7 AA402352	64.30	16.50	17.00	3.90	3.78	+	+	+	+
21400		4.67	1.55	1.00	3.02	4.66	+	+	+	+
	2 AA777488	42.54	7.72	19.71	5.51	2.16	+	+	· <u>-</u>	•
15026		13.66	3.31	3.87	4.12	3.53	+	+	-	+
28989		7.82	1.84	2.30	4.25	3.40	+	+	+	+
	1 AA047441	10.48	3.68	2.19	2.85	4.78	-	+	+	+
	5 AA449837	7.66	2.78	1.58	2.76	4.86	_	+	+	+
20808		7.90	1.98	2.19	3.99	3.60	+	+	+	+
74550	3 AA625995	42.62	11.14	11.32	3.83	3.77	+	+	+	+
509 <b>73</b>	1 AA045699	39.41	23.94	6.63	1.65	5.95	-	-	+	+
	3 AA136386	3.93	2.97	0.63	1.32	6.27	-		+	+
	7 AA054701	39.47	7.39	17.54	5.34	2.25	+	+	-	-
	16 AA461467	6.21	4.35	1.01	1.43	6.16		-	+	+
81423	S AA465598	12.37	6.41	2.19	1.93	5.65		•	+	+
							•			

				Ta	able 1-3					
53022	R15880	1.34	0.70	0.24	1.90	5.63	•	•	+	+
377246 A		12.60	3.02			3.34	+	+	+	-
814662 A		1.12	0.67			5.84	•	_	+	+
45327	H08548	8.18	2.30			3.94	•	+	+	+
51700	H22853	3.66	1.21			4.47	-	+	+	+
80050	T63893	8.15	3.84	1.52	2.12	5.35	-	-	+	+
	AA172372	14.61	10.98	2.38	1.33	6.14	-	-	+	+
•	AA460346	5.39	1.25	1.71	4.31	3.15	+	+	+	+
	4A490611	0.59	0.37	0.10	1.58	5.87	-	-	+	+
	AA100674	225.66	92.92	45.17	2.43	5.00	+	-	+	+
	AA496957	9.40	3.54	1.97	2.65	4.76	+	+	+	+
	AA432075	17.23	6.93	3.50	2.49	4.93	-	•	+ .	+
	AA918370	401.67	285.67	66.91	1.41	6.00	-	-	+	+
	AA988746	9.00	4.61	1.65	1.95	5.45	-	-	+	+
	AA996028	23.76	9.09	4.98	2.61	4.77	•	+	+	+
	AA495819	14.29	3.54	4.28	4.03	3.34	+	+	-	+
	AA460675	26.07	23.22	4.18	1.12	6.24	-	•	+	+
49302	H15675	4.88	0.86	2.90	5.68	1.68	+	+	-	-
	AA436142	12.77	7.46	2.26	1.71	5.64	•	-	+	+
207029		81.88	14.79	45 41	5.54	1.80	+	+	-	•
	AA458747	1.34	1.36	0.21	0.99	6.35	-	-	+	+
	AI015679	3.58	1.45	0.74	2.47	4.85	-	-	+	+
	AA460719	0.66	0.83	0.10	0.80	6.50	-	-	+	+
238821		6.54	2.22	1.50	2.94	4.35	+	+	+	+
129024	1	4.30	1.04	1.36	4.13	3.16	+	+	+	+
	AA430718	16.14	6.63	3.34	2.43	4.83	-	-	+	+
28907		37.48	15.32	7.78	2.45	4.82	+	-	+	+
163728	4	4.01	2.58	0.70	1.55	5.71	-	-	+	+
26369	1	4.23	0.93	1.58	4.57	2.68	+	+	-	+
	7 AA683050	35.22	12.06	8.15	2.92	4.32	+	+	+	+
3238			1.18	0.41	1.85	5.38	-	+	+	+
	0 AA130677		1.75	0.31	1.10	6.12	-	-	+	+
	6 AA465148		0.58	0.37	2.81	4.40	-	+	+	+
	35 AA074258		3.46	3.22	3.48	3.73	+	+	+	+
41631	<b>1</b>		2.66	0.81	1.69	5.52	-	-	+	+
	66 AA448402		3.84	2.97	3.14	4.05		+	+	+
2136			4.10	3.27	3.19	3.99		+	+	+
	07 AA88471	3 3.68		1.07	3.72	3.44		+	+	+
3756	82 AA03222			6.60	2.45	4.70		-	+	+
5105	76 AA05588			27.52	2.34	4.81		-	-	+
2429					3.86	3.28		+	-	+
8781	27 AA77542	2 3.53			4.07	3.05		+	+	+
8539	968 AI79117	3 12.10			4.23	2.87		4	+	+
287					3.97	3.10			+	+
	927 AA48644				1.18	5.83		+	+	+
	042 AA67676				3.63	3.4		_	+	+
	281 AA6702				2.06			-	+	+
	701 AA0111				2.25			+	+	+
	750 N362				3.40			+	•	+
	819 AA4903				3.41			-	+	+
	7807 AA5985							+	•	-
	5749 AA6639							+	-	•
	4895 N211						86 +	+	•	-
82.	5207 AA5041	120 11.	96 2.9	92 4.19	4.05	, 2.0	~~ I			

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					Table 1-3					
461476	AA705041	0.78	0.61	0.14	1.28	5.64	•	-	+	+
323474	W45623	1.25	0.32	0.41	3.87	3.06	+	+	+	+
744846	AA625675	2.69	0.70	0.87	3.83	3.10	+	+	-	+
741474	AA401111	4.93	1.40	1.45	3.51	3.40	+	+	+	+
	AA709044	12.93	4.03	3.49	3.21	3.70	-	+	+	+
	AA777705	3.81	0.83	1.64	4.58	2.32	+	+	+	-
878544	AA775863	19.20	6.78	4.73	2.83	4.06	+	+	+	+
755239	AA422139	17.93	9.29	3.62	1.93	4.95	-	-	+	+
259973	N47829	1.66	1.95	0.28	0.85	5.99			+	+
290370	N62301	0.62	0.48	0.11	1.30	5.53	-	-	+	+
	AA425534	15.94	8.17	3.27	1.95	4.87	_	-	+	+
	AA406505	7.78	2.14	2.45	3.63	3.18	-	+	•	+
290841	N99711	6.35	2.89	1.38	2.20	4.61	•	+	+	+
50606	H17763	14.94	6.00	3.47	2.49	4.31	-	-	+	+
	AA873159	5.15	1.21	2.03	4.26	2.53	+	+	+	-
252314	H87175	0.69	0.62	0.12	1.11	5.66	_	-	+	+
	AA455940	1.75	0.86	0.37	2.04	4.73		_	+	+
	AA620556	18.86	7.00	4.63	2.69	4.07	+	-	+	+
	AA677165	20.56	33.95	3.34	0.61	6.16		_	+	+
	AA699427	7.29	2.43	1.95	3.00	3.74	+	+	+	+
	AA976714	6.44	3.55	1.31	1.81	4.92	_	-	+	+
	AA150502	0.81	0.59	0.15	1.36	5.37	-	-	+	+
	AA133044	559.22	179.15	155.61	3.12	3.59	+	+	+	+
245547		5.86	1.63	1.88	3.59	3.12	+	+	+	+
200599		13.42	5.61	3.11	2.39	4.32	-	+	+	+
139771	ı	7.72	6.17	1.42	1.25	5.44	_	-	+	+
	AA705376	8.32	6.33	1.55	1.31	5.38	_	-	+	+
	AA464580	4.37	1.58	1.11	2.76	3.92	-	+	+	+
279482	•	8.66	3.25	2.16	2.67	4.00	+	-	+	+
127396	l .	14.07	5.12	3.61	2.75	3.90	-	+	+	+
	AA490609	4.64	2.75	0.94	1.69	4.95	-	-	+	+
	AA609323	38.01	9.06	15.59	4.20	2.44	+	+	+	-
	AA287375	9.94	3.00	3.00	3.31	3.31	-	+	-	+
250869	1	123.92	32.54	44.10	3.81	2.81	+	_	+	•
	AI025120	5.09	1.33	1.84	3.84	2.77	+	+	-	+
	AA779391	18.03	7.40	4.35	2.44	4.14	-	+	+	+
	AA609368	7.90	4.94	1.59	1.60	4.98	-	-	+	+
81578		12.77	4.82	3.25	2.65	3.93	-	+	+	+
	4 AA434360	103.13	30.63	32.18	3.37	3.20	+	+	+	+
	5 AA621761	14.86	3.64	5.98	4.08	2.48	+	+	-	-
	6 AA459100	18.28	10.68	3.77	1.71	4.85	-	-	+	+
7791		14.99	17.48	2.63	0.86	5.70	-	•	+	+
26011		0.88	0.86	0.16	1.03	5.52	-	-	+	+
	3 AA458551	17.84	5.98	5.02	2.98	3.55	-	+	-	+
21040		34.02	7.84	15.49	4.34	2.20	+	+	•	-
12482		30.85	11.28	8.13	2.73	3.79	+	•	+	+
	9 AA169379	17.74	7.51	4.26	2.36	4.17	+	+	+	+
	3 AA278473	4.77	2.81	0.99	1.70	4.82	-	-	+	+
	3 AA058818	15.01	6.81	3.48	2.20	4.32	•	•	+	+
19160		22.14	4.64	12.72	4.77	1.74	+	+	-	-
	5 AA012866	40.29	13.93	11.17	2.89	3.61		+	+	+
	7 AA491265	0.74	0.41	0.16	1.81	4.68	-	-	+	+
	71 AA 181179	5.74	2.03	1.58	2.82	3.64	+	+	+	+
150000	00 AA885642	6.11	3.34	1.32	1.83	4.63	-	-	+	+

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				T	able 1-3						
813073	AA455689	3.24	0.88	1.18	3.70	2.75	•		+	-	•
	AA504536				1.48	4.97			-	+	+
	AA159497	0.46			2.95	3.50	+		-	+	+
	4A457681	1.90			0.74	5.71	-		-	+	+
		13.36		3.37	2.48	3.97	_		-	+	+
	AA644080			7.5 <b>8</b>	2.10	4.34	-		-	+	+
and the second second	AA465614			0.96	2.29	4.14	_		_	+	+
	AA989473	3.96		1.61	2.57	3.86	+		+	+	+
	AA629117	6.21	2.41		2.57	3.86	_	-	•	+	+
	AA488784	2.63		0.68	3.02	3.40		-	+	_	+
290642	N99659	7.76	2.56	2.28		3.13			+	_	+
121489	T97303			11.09	3.29	4.82		- -	_	+	+
	AA489647	7.42	4.65	1.54	1.59	3.58			+		+
86160	T72336	2.10	0.74	0.59	2.82			-	· -	+	+
	AA862465	41.61	40.32	7.76	1.03	5.36		-	+	_	
	AA598470	2.07	0.49	0.98	4.25	2.12		+	+	_	_
	AA488238	67.93		36.21	4.50	1.88		+		+	+
504544	AA149096	12.31	4.73	3.28	2.60	3.76		+	-	+	+
1466631	AA883679	1.21	0.71	0.26	1.71	4.63	ı	•	-		+
898109	AA598483	6.77	3.15	1.62	2.15	4.19		-	-	+	+
280763	N50563	0.53	0.50	0.10	1.05	5.28		-	-	+	+
	AA875936	13.96	4.83	4.08	2.89	3.43	1	-	+	•	
194364		5.44	1.19	3.10	4.56	1.76		+ .	+	-	<del>-</del>
	AA699895	3.24	0.96	1.10	3.36	2.96	1	•	+	<del>-</del>	+
	AA430351	41.57	23.92	9.10	1.74	4.57	1	-		+	+
2906	•	3.34	1.66	0.78	2.01	4.29	1		-	+	+
	4 AA171459	1.70	1.73	0.32	0.98	5.32	1	-	-	+	+
2661		4.62	2.68	1.01	1.73	4.56	1	-	-	+	+
	9 AA045115	4.41	1.79	1.16	2.47	3.82		+	-	+	+
163184		25.01	12.24	5.91	2.04	4.24	1	-	-	+	+
20411	1	3.38	0.87	1.40	3.86	2.42	1	+	+	-	-
	4 AA432312	9.75	3.24	2.99	3.01	3.26		+	-	+	•
	34 AA137144	6.35	1.83	2.27	3.47	2.80	) [	+	+	+	+
	88 AA486145	23.03	5.46	11.26	4.22	2.05	; [	+	+	-	-
	58 AA082943	9.57	4.00	2.48	2.39	3.86	5	-	-	•	+
	53 AA446906	11.76	3.47	4.12	3. <b>39</b>	2.85	5	-	+	-	+
1961		11.19	4.82	2.86	2.32	3.9	1	-	-	-	+
475	1	8.07	1.88	4.18	4.30	1.93	3	+	+	-	-
473		4.22	1.17	1.61	3.61	2.63	2	-	+	-	•
	37 AA491249		31.45	19.54	2.38	3.8	3	•	-	+	+
1947			3.86	1.07	1.35	4.8	6	•	-	+	+
1503			2.31	1.35	2.29	3.9	2	•	-	+	+
	784 AA486200		118.59	94.88	2.75	3.4		+	+	+	+
	286 H12190		0.61	0.99	3.83	2.3		-	+	-	-
	771 AA677287		0.70	1.13	3.82	2.3		+	+	-	-
	208 R38412		3.51	1.85	2.13	4.0		-	-	+	+
			3.09	2.81	2.95	3.2		-	+	-	+
	1		0.96	0.19	1.01		16	-	-	+	. +
	419 N2497 410 AA41225		0.58	0.19	0.92		24	-	-	+	+
			1.57	1.81	3.30		85	+	+,	-	+
	973 AA97729				2.36		77	-		-	+
	668 AA50506				2.25		.88	-	-	+	+
	2864 N2443				1.52		.62			+	+
	5201 AA62614						.70		•	+	+
	9450 A A 68024						.18	+	•	+	-
3'	7980 R613	72 0.84	0.29	0.27	2.50	, ,					

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					Table 1-3					
124252	R02439	6.06	4.74	1.25	1.28	4.85	-		+	+
238661	H81543	2.74	2.18	0.56	1.26	4.86	•	-	+	+
593658	AA160080	1.08	1.02	0.21	1.07	5.04	•	_	+	+
785616	AA449004	11.47	3.12	4.70	3.67	2.44	-	+		-
84022	T70901	6.97	2.71	1.97	2.57	3.54	•	_	+	+
511632	AA126951	7.85	2.71	2.46	2.89	3.20	•	+	-	+
34134	R44734	2.73	1.33	0.68	2.06	4.03	•	-	+	+
39885	R52543	0.94	0.30	0.31	3.11	2.98	-	+	-	+
	AA488014	21.70	7.54	6.78	2.88	3.20	+	-	+	-
49710	H28973	3.21	1.47	0.82	2.18	3.89	-	-	-	+
42803	R60014	2.96	0.81	1.23	3.66	2.41	+	+	+	•
134969	R31789	6.81	4.59	1.48	1.48	4.58	•	-	+	+
75415	T57609	51.71	24.46	13.09	2.11	3.95	-	~	•	+
	AA488626	49.18	13.00	21.59	3.78	2.28	•	+	-	-
137189	R36409	5.26	1.96	1.56	2.68	3.37	-	+	-	+
45578	H08016	5.05	1.63	1.72	3.10	2.94	-	+	-	+
	AA680186	5.55	1.53	2.30	3.62	2.41	+	-	+	-
67654	T49539	15.79	6.24	4.50	2.53	3.51	+	+	+	+
	AA102591 AA459765	78.20 2.34	22.40 1.80	30.93 0.50	3.49 1.30	2.53 4.72	-	-	+	+
1	AI005519	2.34 25.52	13.27	6.23	1.92	4.09	-	-	+	+
	AA083485	12.81	2.43	17.02	5.26	0.75	+	+	-	_
	AA489696	0.49	0.34	0.11	1.42	4.58	<u>.</u>		+	+
	AA425692	21.34	15.28	4.63	1.40	4.61	-	-	+	+
	AA679569	4.23	2.23	1.03	1.89	4.10		-	+	+
	AA481256	9.59	4.44	2.50	2.16	3.84	-	-	+	+
196650		10.93	6.59	2.52	1.66	4.33	•	-	+	+
	AA398521	26.71	9.34	8.54	2.86	3.13	•	+	-	+
	AA872436	2.96	0.83	1.23	3.58	2.40	+	+	-	-
	AA485748	6.36	4.30	1.41	1.48	4.50	-	-	+	+
366132	AA062805	6.22	1.68	2.75	3.72	2.26	-	+	-	-
	AA287067	16.03	15.09	3.27	1.06	4.91		-	+	+
	AA442301	12.77	6.52	3.19	1.96	4.01	-	-	-	+
	AA788970	7.60	2.71	2.40	2.80	3.17	+	+	+	+
210862	1	2.56	1.16	0.68	2.21	3.76	-	-	+	+
	AA083671	8.46	7.83	1.73	1.08	4.88	-	-	+	+
	AA425773	21.90	19.94	4.51	1.10	4.86	-	•	+	+
	AA405597	1.03	1.35	0.20	0.76	5.19	•	-	+	+
66902	AA460251	17.83 3.21	7.09	5.20 1.03	2.51 2.82	3.43 3.12	+	+	<b>-</b>	+
	T67444 B AA505162	4.03	1.14 1.32	1.03	3.06	2.88	_ <del>_</del>	+	-	+
20128		6.42	4.16	1.46	1.55	4.39		· -	+	+
	2 AA987644	1.97	0.81	0.57	2.45	3.48	+	_	+	+
	6 AA018979	4.39	2.04	1.16	2.15	3.77	_	-	+	+
	1 AA864875	12.82	5.49	3.58	2.33	3.58		-	-	+
	5 AA599717	3.26	1.43	0.90	2.28	3.63		+	+	+
10915		5.46	1.70	2.03	3.20	2.69	<b>.</b>	+	-	-
5076		0.43	0.23	0.11	1.89	4.00	-	-	+	+
35699		21.95	7.93	7.04	2.77	3.12	-	+	-	+
	4 AA996104	6.06	2.17	1.97	2.79	3.07	+	-	+	+
	3 AA488865	4.03	2.24	0.99	1.80	4.06		-	+	+
25946		3.69	1.65	1.02	2.24	3.62	+	-	+	-
	8 AA425205	26.40	17.78	6.05	1.49	4.37		•	+	+
62699	00 AA 190850	7.39	3.82	1.89	1.93	3.91	-	-	+	+

				•	Table 1-3		_			
7437491	AA634291	5.10	1.65	1.86	3.10	2.74	-	+	•	-
79000	T62164	6.24	3.53	1.54	1.77	4.05	•	-	+	+
23576	T77385	3.58	1.59	1.00	2.25	3.57		-	-	+
	AA029490	71.61		36.97	3.87	1.94	+	-	+	-
	AA421256	0.85	1.00	0.17	0.85	4.96	-	-	+	+
415437	W81124	3.52	1.02	1.50	3.46	2.35	-	+	-	-
197779	R93515	8.61	6.61	1.91	1.30	4.50	-	-	+	+
	AA902831	0.85	0.55	0.20	1.55	4.25	_	-	+	+
	AA126009	24.31	24.09	5.08	1.01	4.79	-	•	+	+
	AA181995	6.17	2.49	1.86	2.47	3.32	-	-	-	+
	AA456616	24.75	12.84	6.40	1.93	3.87	_	•	+	+
	AA015658	28.16	15.79	7.03	1.78	4.01	_	-	+	+
	AA682627	13.81	4.59	4.98	3.01	2.77	•	+	•	+
	AA280288	0.50	0.30	0.12	1.68	4.10	-	-	+	+
296788		3.27	0.67	3.50	4.85	0.93	+	+	-	-
1638852	•	1.21	0.50	0.36	2.41	3.36	-	-	-	+
1636868		5.07	2.89	1.26	1.76	4.01	-	-	+	+
	AA705142	11.87	12.89	2.45	0.92	4.84	-	_	+	+ ·
357364	1	0.72	0.27	0.23	2.67	3.09	_	_	+	+
	7 AA169645	24.81	12.50	6.57	1.98	3.78	_	_	+	+
	AA486412	2.20	1.38	0.53	1.60	4.16	-	-	+	+
111812	L L	5.56	1.89	1.98	2.94	2.81	-	+	•	+
	6 AA443969	5.25	4.09	1.17	1.28	4.47	-	-	+	+
29204:		9.39	2.67	4.19	3.51	2.24	-	+	~	-
	8 AA634379	3.34	1.15	1.17	2.90	2.85	+	+	+	+
	7 AA629910	6.68	2.72	2.04	2.46	3.28	_	-	-	+
	4 AA634360	5.09	1.90	1.66	2.68	3.06	_	-	-	+
	9 AA019445	3.03	1.24	0.92	2.44	3.30		-	+	+
	0 AA127100	4.09	1.46	1.39	2.80	2.93	-	+	-	+
	8 AA630084	6.21	2.55	1.89	2.44	3.29	+	-	+	-
	35 AA189113	15.94	8.27	4.20	1.93	3.79	-	-	+	+
	22 AA021586	4.15	1.53	1.39	2.72	3.00		-	+	+
26446		0.77	0.84	0.16	0.91	4.79	-	-	+	+
2326		4.02	1.27	1.58	3.16	2.55	+	+	-	+
	10 AA599064	3.18	0.89	1.49	3.56	2.14		-	-	-
	61 AA457767	2.59	2.11	0.58	1.23	4.47		-	+	+
	62 AA400732	2.81	1.76	0.69	1.60	4.10	-	-	+	+
	11 AA169192	34.63	11.27	13.23	3.07	2.62		+		+
2513	1	9.21	3.78	2.83	2.44	3.25	;   -	+	-	+
	88 AA497127	7.72	2.72	2.70	2.83	2.85	5   -	+	-	+
	48 AA017383		1.69	0.82	1.85	3.83	3   -	-	+	+
	02 AA521409		1.51	1.45	2.78	2.89	+	-	+	•
	32 AA625628		13.65	18.48	3.25	2.4	0   -	+	•	-
	447 AA630800		2.09	6.13	4.22	1.4	4 +	+	•	•
	715 R44078			0.71	3.73	1.9	3   -	+	-	-
	549 AA456827				1.17	4.4	9 -	-	+	+
	443 AA49593				1.73	3.9	2 -	-	+	+
	602 AA86468				2.40	3.2	5 -	-	•	+
	103 AA62636				2.67	2.9		+	-	+
	258 AA44701					2.0	ю +	+	•	-
	836 AA60867				2.56			-	+	-
	052 AA17660						53 -	-	+	+
	7343 A101535							-	+	+
	2325 AA70551					3.	33   +	•	+	•
	•									

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					Table 1-3					
878284	AA670280	4.41	2.26	1.20	1.95	3.66	-		+	+
	AA042812	7.64	5.06	1.86	1.51	4.10	•	_	+	+
	AA670296	5.34	1.86	1.95	2.87	2.73	•	+	•	+
251019	H97778	45.62	22.03	12.95	2.07	3.52	-	•	+	+
258167	N40582	17.24	8.15	4.96	2.12	3.48	-	-	+	+
416833	W87312	6.07	2.16	2.18	2.81	2.78		+	-	+
186768	H51914	9.43	5.75	2.39	1.64	3.95	-	-	+	-
	AA976063	8.20	5.56	1.99	1.47	4.11	-	-	+	+
49728	H29198	1.27	0.57	0.38	2.22	3.36	-	-	_	+
125722		3.28	1.21	1.14	2.72	2.87	-	+	<u>-</u>	+
613237	AA181767	1.07	0.78	0.25	1.37	4.22	-	-	+	+
526945	AA112515	1.14	0.40	0.41	2.82	2.77	-	+	-	+
79032	T61960	37.58	31.85	8.53	1.18	4.40	-	-	+	+
843139	AA485922	39.96	18.13	11.84	2.20	3.37	•	-	-	+
841302	AA487213	7.41	3.64	2.09	2.04	3.54	+	-	+	-
191978	H38623	47.95	29.72	12.11	1.61	3.96	-	-	+	+
135900	R33609	9.44	5.00	2.57	1.89	3.68	-	•	+	+
713109	AA282983	2.44	0.98	0.79	2.48	3.08	-	-	-	+
626640	AA191512	8.33	4.14	2.35	2.01	3.54	•	-	+	+
322914	W45148	9.34	3.68	3.10	2.54	3.02	-	-	-	+
251517	H96605	0.73	0.80	0.16	0.91	4.64	•	•	+	+
840878	AA482228	25.81	27.85	5.59	0.93	4.61	-	-	+	+
626716	AA191245	21.60	13.98	5.41	1.55	3.99	•	•	+	+
291345	W02978	70.40	31.69	21.23	2.22	3.32	+	-	+	+
805 <b>0</b> 0	1	140.81	52.48	49.33	2.68	2.85	•	+	+	+
	AA485201	3.64	2.20	0.94	1.65	3.88	-	-	+	+
	AA703079	168.72	131.76	39.68	1.28	4.25	-	-	+	+
	AA705077	0.62	0.38	0.16	1.66	3.87	-	-	+	+
	8 AA001234	4.05	3.19	0.95	1.27	4.25	-	-	+	+
13763		5.06	1.71	1.97	2.95	2.57	-	+	•	+
	4 AA664101	17.75	8.17	5.31	2.17	3.34	+	-	+	-
10931		12.35	4.01	5.08	3.08	2.43	+	+	+	•
	8 AA399198	4.92	2.96	1.28	1.66	3.85	•	-	+	+
	8 AA708342	34.95	10.61	15.79	3.29	2.21	•	+	-	-
12070		19.60	12.65	4.95	1.55	3.96	-	-	+	+
	0 AA486723 4 AA994757	1.46 21.83	1.23 14.39	0.34 5.47	1.18 1.52	4.32	•	-	- +	+
	8 AA418755	5.70	14.39	2.18	2.88	3.99 2.61	•	+		+
25143		0.73	0.90	0.16	0.81	4.68	•	-	+	+
7504		3.46	0.93	1.96	3.72	1.77	_	+		`
	1 AA405984	1.89	1.02	0.52	1.85	3.63	-		+	+
	6 AA133165	3.48	1.03	1.66	3.38	2.10	-	+		•
34458		53.28	21.92	17.55	2.43	3.04		+	_	+
25067		2.66	1.12	0.86	2.37	3.09	+	-	+	_
	0 AA449644	4.01	1.01	2.72	3.98	1.48	+	+	•	-
	29 AA489670	1.94	1.47	0.47	1.32	4.13		-	+	+
	0 AA826237	13.40	6.84	3.84	1.96	3.49	•	-	•	+
	18 AA010608	10.87	4.02	3.96	2.71	2.74	_	+	-	+
455		1.66	0.76	0.51	2.17	3.28	-	-	+	+
2519		7.48	2.74	2.76	2.73	2.71	-	+	•	-
2533		1.52	1.34	0.35	1.14	4.30		-	+	+
	32 AA121428	2.68	0.85	1.17	3.15	2.28	-	+	-	-
	44 AA630545	15.78	6.78	5.09	2.33	3.10	+	+	+	+
8982	42 AA598621	10.34	7.74	2.53	1.34	4.09		-	+	+

				τ	able 1-3			_			
395625 AA	7576D4 1	14.45	6.11	4.73	2.37	3.06		-		-	+
767406 AA				0.10	0.98	4.45	-	-		+	+
				4.07	2.17	3.25	-	-	•	+	+
1031598 AA				3.37	3.88	1.54	+	+		-	-
785913 A				2.75	4.19	1.22	+	+		-	-
1				2.59	0.95	4.46	-	-		+	+
141505	R72977	6.38	1.86	3.21	3.42	1.99	-	+		-	-
298155	N70794	2.85	1.29	0.89	2.21	3.19	-	-		•	+
745283 A		7.94	5.76	1.97	1.38	4.02	-	-		-	+
26230	R12449	5.08	2.29	1.60	2.22	3.18	-				+
743851 A		8.22	3.22	2.89	2.55	2.84	-	+		-	+
884531 A		7.71	4.73	2.05	1.63	3.76	-	-		+	+
345032	W76319	20.78	9.11	6.69	2.28	3.11	-			•	+
i i		4.95	3.66	1.23	1.35	4.03	-	-		+	+
729953 A 815017 A		14.81	9.71	3.84	1.52	3.85	-			-	+
1		9.26	4.85	2.67	1.91	3.47	-	-		-	+
308466	N95495	1.21	1.11	0.28	1.09	4.29	-	_		+	+
773345 A		4.93	3.60	1.23	1.37	4.00	_	-	,	+	+
811766 A	N59119	4.93	1.38	2.37	3.40	1.97	-	4	<del>.</del>	-	•
246869		12.44	4.26	5.09	2.92	2.44	+	4	<b>-</b>	-	-
1473300 A		10.98	6.11	3.08	1.80	3.56	_			-	+
	A702247	6.44	3.83	1.75	1.68	3.68	-		-	-	+
	AA446643		5.84	4.25	2.26	3.10	٠.		+	+	+
293243	N68686	13. <b>1</b> 7 5. <b>8</b> 6	3.56	1.58	1.65	3.71	_		-	-	+
	AA598996		2.43	3.67	3.22	2.13			+	•	-
	AA424706	7.81 2.98	2.43	0.76	1.43	3.93	-		-	+	+
299723	N75055	2.26	2.09	0.73	1.08	4.27	-		-	+	+
I I	AA705072	1.21	0.48	0.33	2.50	2.85	1 .		+	•	+
82905	T69305	1.74	1.05	0.47	1.65	3.69			-	-	+
	AA489121 T97457	6.37	6.46	1.46	0.99	4.35			•	+	+
121436	AA599078	7.48	3.82	2.21	1.96	3.38			-	-	+
	AA456331	4.72	2.68	1.32	1.76	3.57			•	+	+
	t e	4.72	3.20	1.10	1.37	3.97	•		-	+	+
32509 346889		0.89	0.21	0.79	4.21	1.12		•	+	-	-
145388		2.66	0.94	1.06	2.83	2.50			+	•	-
	AA682631	1.85	1.07	0.51	1.73	3.59		•	-	+	+
46411	1	3.07	2.00	0.81	1.54	3.7			-	+	+
	AA399269	3.50	2.68	0.87	1.30	4.0			-	+	+
	AA490490	1.57	1.72	0.36	0.91	4.4		-	-	-	+
288695		3.41	0.88	2.36	3.86	1.4	4	+	+		-
	2 AA436595	7.06	3.62	2.10	1.95	3.3	6	-	-	•	+
	5 AA773478	5.09	3.00	1.41	1.70	3.6		-	-	-	+
13313		11.12	4.91	3.67	2.27	3.0		•	-	+	+
	1 AA452955	12.11	3.01	9.65	4.03	1.2		+	+	-	-
41786			9.26	3.54	1.46	3.8	32	-	-	•	+
	O AA173109		6.62	1.96	1.21		07	-	-	+	+
	6 AA934762		5.52	2.06	1.43		84	-	•	-	+
35845			23.51	10.38	1.62		66	-	-	-	+
	0 AA464531		20.86	7.03			94	-	-	•	+
	22 AA001403		5.49				59	-	-	-	+
5192			1.72				.67	-	-	+	-
	94 AA41882		1.83				.93	+	•	+	•
	92 AA47913						.04	-	-	-	+
2543							.92	+	•	+	+
2343	101 120112	. 10.20					•				

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					Table 1-3		_	_		
1635110 A		38.13	17.09	12.62	2.23	3.02	•	•	-	+
121997	T98253	5.10	1.52	2.70	3.36	1.89	+	+	-	-
51011	H19242	4.64	2.46	1.38	1.88	3.36	•	-	+	+
283034	N51280	9.61	3.05	4.60	3.15	2.09	+	+	-	-
564537 A	A 12 1732	1.79	1.93	0.42	0.93	4.31		-	+	+
82734	T73651	7.21	6.90	1.72	1.05	4.18	_	_	+	+
1631682 A		1.61	0.53	0.74	3.03	2.19	-	+		-
290378	N64508	17.10	17.02	4.07	1.00	4.20	-	-	+	+
262865	H99650	30.48	11.78	11.64	2.59	2.62		+	_	+
1031698 A		6.26	3.13	1.95	2.00	3.21	•	-	-	+
109200	T81107	3.17	0.93	1.78	3.42	1.79	+	+	•	· •
824358 A		0.96	0.68	0.25	1.42	3.78	•	•	+	+
32684	R20424	24.32	17.98	6.32	1.35	3.85			+	+
240430	H78097	3.52	1.28	1.43	2.74	2.46	+	_	-	
1475200		28.16	22.56	7.13	1.25	3.95		-	+	+
	AA421073	5.02	2.47	1.59	2.03	3.16	-	_	· _	+
245860	N55339	30.36	12.47	11.02	2.43	2.76		+	+	+
	AA630346	2.22	0.68	1.15	3.26	1.92	+	+	· -	•
	AA490802	1.03	0.63	0.29	1.63	3.55		<u>-</u>	-	+
1435029		1.84	0.67	0.75	2.74	2.44	_	+		_
	AA620446	2.43	1.63	0.66	1.49	3.69	_	<u>.</u>	+	+
22731	R45264	4.46	2.73	1.26	1.64	3.54	-	_	· +	+
	AA447742	34.08	17.95	10.41	1.90	3.27	_	•	+	·
258521	N40743	3.33	1.39	1.20	2.40	2.77	+	_	, +	_
	AA988798	135.32	117.37	33.70	1.15	4.01	_	_		+
144926	R78559	9.20	3.96	3.24	2.32	2.84	-	+	<u>-</u>	+
	AA598758	15.87	13.03	4.03	1.22	3.94	_	•	+	+
	AA478268	7.20	4.16	2.10	1.73	3.42	_	_	+	+
	AA877347	8.03	3.31	2.94	2.43	2.73	_	_		+
277660	N49405	14.16	5.37	5.65	2.64	2.51	_	+	-	+
	AA486261	17.59	8.78	5.60	2.00	3.14	_	<u>.</u>	_	+
	AA426227	5.89	3.00	1.85	1.96	3.18	_	_	_	+
	AA456721	22.70	10.74	7.51	2.11	3.02	_	_	_	+
	AA262573	3.36	1.79	1.03	1.88	3.26	•	_	_	+
	AA130042	18.15	5.54	9.79	3.28	1.85	+	+		-
1631762	AI025126	46.32	36.08	12.06	1.28	3.84			_	+
343167	W67309	15.62	7.28	5.24	2.15	2.98	_	_	_	+
30793	R18258	7.89	2.84	3.37	2.78	2.34	_	+	_	_
82874	T69348	5.14	2.70	1.60	1.90	3.21		-	-	+
209167	H63668	3.29	1.15	1.46	2.86	2.25	+	+	+	_
	AA443177	9.91	4.54	3.39	2.18	2.92		-	-	+
	AA464700	3.85	1.37	1.68	2.81	2.30	-	+	•	
249705		62.65	37.20	18.34	1.68	3.42		-	-	+
178137	1	238.41	190.87	61.93	1.25	3.85		-	+	+
	AA485216	1.22	1.44	0.29	0.84	4.25	_	_	+	+
294259		4.21	2.21	1.32	1.90	3.19	l .	-	+	+
248454		3.70	1.95	1.16	1.89	3.19	١.	_	+	
	AA151796	50.88	17.89	22.72	2.84	2.24	+	_	+	
	AA416833	4.45	2.31	1.41	1.93	3.15	1	•	+	+
259350		22.84	6.64	13.93	3.44	1.64		+	•	-
	AA894687	3.57	1.31	1.53	2.73	2.34		+	-	+
	AA195601	5.48	2.72	1.79	2.02	3.06		· -	•	+
	AA056001	2.59	1.39	0.81	1.87	3.20	+	_	+	_
	AA458578	10.44	6.85	2.94	1.52	3.55	I	-	+	+
-2.,,	,		05	2.77		ر د. د	1 '		•	

		1		٦	Table 1-3		_			
1500324 A	A885 <b>8</b> 19	6.77	4.55	1.89	1.49	3.58	•	-	+	+
167205	R90934	7.85		3.93	3.06	2.00	+	+	•	-
594693 A		7.33		2.44	2.05	3.00	-	-	+	+
868119 A		10.00	4.58	3.49	2.18	2.87	-	-	-	+
502463 A		5.77	2.99	1.85	1.93	3.12	-	-	•	+
342271	W61215	3.90	1.09	2.69	3.60	1.45	+	+	-	•
256680	H96392	1.72	1.47	0.44	1.17	3.88	-	-	•	+
430179 A	A010091	51.95	32.22	15.13	1.61	3.43	-	-	+	-
897153 A	A676961	10.77	5.36	3.55	2.01	3.04	-	-	•	+
25517	R11888	8.52	4.10	2.87	2.08	2.96	•	-	•	+
1632141	A1004916	2.15	0.93	0.79	2.31	2.74	-	-	•	+
814018 A	A455714	0.98	0.77	0.26	1.28	3.77	-	-	+	+
843234 A	A488588	7.02	2.32	3.49	3.03	2.01	-	+ ,	-	-
823578 A	A497118	0.56	0.53	0.14	1.06	3.98	-	-	•	+
42636	R61780	1.11	0.32	0.72	3.50	1.54	-	+	•	•
784830	\A448289	1.83	0.63	0.85	2.88	2.15	-	+	-	-
244652	N52911	24.70	12.93	7.92	1.91	3.12	-	-	+	-
950709	<b>4A608583</b>	4.17	1.22	2.62	3.43	1.59	+	+	-	-
1486260	AA922326	13.23	7.43	4.08	1.78	3.24	-	-	+ '	+
878152	AA775431	4.20	3.49	1.10	1.20	3.82	-	<del>-</del>	-	+
130773	R22156	3.71	1.21	1.90	3.07	1.95	-	+	•	•
1	AA129171	2.32	0.70	1.34	3.29	1.73	-	+	-	+
	AA126356	33.96	23.00	9.58	1.48	3.54	-	-	+	+
295206	N75947	3.16	1.64	1.02	1.92	3.10	-	-	Ŧ	_
281072	N50912	. 5.02	1.92	2.10	2.62	2.39	+	•	•	-
	AA482198	2.04	0.70	0.98	2.91	2.09	-	+	-	+
143306	R74169	10.42	4.91	3.62	2.12	2.88	-	- +	-	· -
247609	N58148	3.15	1.20	1.33	2.63	2.37	-	<b>T</b>	-	+
250678	1	4.24	1.94	1.51	2.18	2.82 3.62	•	-	+	+
	AA458943	0.47	0.34	0.13	1.38 2.21	2.79	+	_	+	_
	AA504284	3.42	1.55	1.23	1.07	3.92	_	_	+	+
197888		4.97	4.63	1.27 4.15	2.75	2.25	+	_	-	-
261472		9.31	3.39	0.51	2.75	2.64	_	+	_	+
137254		1.35 6.16	0.58 4.38	1.72	1.41	3.58	_	-	+	+
52730		11.64	8.74	3.19	1.33	3.65	} _	-	+	+
76252	T59873 AA598487	4.77	1.81	2.03	2.63	2.35	<b>i</b> -	+	-	•
	1 AA485214	14.69	7.77	4.77	1.89	3.08	-	-	+	+
	2 AA262559	3.59	3.70	0.90	0.97	3.99		-	+	+
23346		14.24	3.73	12.38	3.81	1.15	+	+	•	-
8048		14.69	14.52	3.72	1.01	3.95	<b>!</b> -	-	+	+
32205		11.58	4.27	5.15	2.71	2.25	-	+	-	-
29915		1.37	0.71	0.45	1.92	3.04		-	-	+
13660		10.58	5.59	3.46	1.89	3.06		-	+	+
	0 AA281504		1.16	0.35	1.14	3.81	-	•	-	+
	AA664105		1.34	1.22	2.36	2.59	)   -	-	+	+
2725			4.17	1.80	1.49	3.45	;   -	-	-	+
833			3.35	1.48	1.51	3.43	3   -	•	-	+
	91 AA933888		3.85	5.05	2.80	2.14		+	•	+
	50 AA496628					3.50		•	+	+
	55 AA459679							+	-	•
1482					1.98			-	+	_
1881				1.23	2.30			•	+	-
	09 AA47848					5 2.7	8 +	-	+	•
	•									

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					Table 1-3					
561971 A	A088886	3.81	3.31	1.01	1.15	3.78		•	-	+
	A608546	3.53	2.28	1.04	1.55	3.39	•	-	+	-
266747	N31312	3.05	1.60	1.01	1.91	3.02	-	•	• .	+
129146	R10947	7.97	4.87	2.42	1.64	3.30	•	-	•	+
41940	R59087	10.51	5.11	3.65	2.05	2.88	•	-	•	+
	A452113	1.46	0.42	1.00	3.46	1.46	+	-	•	-
	<b>A</b> 070226	46.55	26.81	14.60	1.74	3.19	•	-	-	+
287528	N62096	1.98	1.06	0.65	1.87	3.05	-	-	+	+
	AA287323	8.70	5.76	2.55	1.51	3.41	-	-	•	+
	AA857632	2.56	1.69	0.75	1.51	3.40	•	•	+	+
248261	N78083	1.51	0.70	0.55	2.15	2.76	-	+	+	+
	AA620458	2.03	1.20	0.63	1.69	3.22	-	-	•	+
123755	R01304	5.06	2.03	2.10	2.49	2.41	+	+ .	+	+
	AA147499	2.26	1.01	0.84	2.23	2.68	-	-	•	+
	AA521247	3.50	2.11	1.08	1.66	3.24	•	-	+	+
1	AA464617	7.62	5.19	2.22	1.47	3.43	-	-	+	+
	AA906997	2.08	0.72	1.04	2.90	2.00	-	+	-	-
416010	W85832	53.71	32.20	16.66	1.67	3.22	-	-	+	-
	AA610071	25.15	13.97	8.15 2.97	1.80	3.09	-	-	+	+
	AA775857	11.10 4.46	9.66 1.54	2.25	1.15 2.90	3.74 1.98	•	- +	•	т
430133	AA010233 W88848	4.40 17.18	1.54	2.23 4.77	1.28	3.60	-	<b>.</b>	+	+
108265	T70810	7.24	5.07	2.10	1.43	3.45	•	•	т	+
	AA625574	29.18	19.75	8.59	1.48	3.40	_		+	, +
	AA025112	4.93	1.78	2.34	2.76	2.11	_	+	-	
112785	T85958	7.75	4.81	2.38	1.61	3.25			-	+
	AA447761	4.96	1.93	2.17	2.58	2.29	_	+	_	-
	AA463932	2.27	1.27	0.74	1.79	3.07	_	-	+	+
	AA399264	10.10	6.03	3.17	1.68	3.18	_	-	-	+
	AA133721	4.59	1.55	2.41	2.96	1.90	+	_	+	-
	AA485018	2.31	1.17	0.81	1.98	2.87	-	-	-	+
50344		0.85	0.39	0.32	2.16	2.69	+	-	+	-
293830	N98283	4.43	1.72	1.96	2.58	2.26	+	-	+	-
213890		14.14	7.99	4.60	1.77	3.07	-	-	+	+
	AA446885	10.62	7.56	3.09	1.41	3.44	-	-	-	+
36 <b>607</b>	•	7.52	6.30	2.06	1.19	3.65	•	-	+	+
	AA278195	4.12	2.45	1.31	1.68	3.15	•	-	-	+
40017		8.29	4.13	2.94	2.01	2.82	-	-	-	+
	AA253444	4.99	2.55	1.74	1.96	2.87	•	-	+	-
	AA122048	5.94	3.29	1.97	1.81	3.02	-	-	-	+
	AA074677	2.82	1.94	0.84	1.46	3.37	-	-	+	+
	AA056484 N48701	7.44 3.18	5.33 1.23	2.17 1.42	1.40 2.59	3.43 2.23	+	-	+	_
74223		5.42	3.10	1.42	1.75	3.07	- T	-	_	+
	AA054643	3.54	1.49	1.46	2.38	2.43	. +	_	+	
	AA281793	1.06	0.58	0.35	1.83	2.98		-		+
	7 AA702480	3.91	1.64	1.61	2.38	2.43		+	_	+
	4 H16851	0.83	0.28	0.44	2.92	1.89	+	+	-	+
	8 AA872323	7.11	4.65	2.16	1.53	3.29	_	-	-	+
	9 AA487552	9.08	4.37	3.33	2.08	2.73	_	-	_	+
	9 14664067	9.00	6.46	2.64	1.39	3.41		•	+	+
	3 AA457737	4.95	1.68	2.67	2.95	1.85	-	+	-	-
5174		3.65	1.79	1.33	2.05	2.75	-	•	+	-
	9 W96280	13.43	4.23	8.34	3.18	1.61	-	+	-	-
	•						•			

					Table 1-3			_		
252453	H87143	20.12	14.53	5.91	1.38	3.40	•	_	+	+
133717	R27767	9.46	2.97	5.91	3.18	1.60	+	+	•	_
	AA449037	55.68	21.05	26.01	2.65	2.14	-	+	•	•
76196	T59668	2.53	0.90	1.29	2.83	1.96	+	+	-	-
322561	W15277	107.98	70.00	33.31	1.54	3.24	-	-	-	+
825833	AA505150	1.95	1.08	0.66	1.81	2.97	-	-	•	+
755765	AA496455	4.01	2.20	1.36	1.83	2.96	-	-	-	+
1416086	AA878210	8.24	4.30	2.88	1.92	2.86	-	-	+	+
358083	W95279	2.45	0.80	1.44	3.07	1.71	-	+	-	-
	AA447731	84.51	49.31	27.59	1.71	3.06	-	-	-	+
	AA133166	· 5.35	3.78	1.60	1.42	3.35	-	•	+	+
	AA455126	27.74	19.39	8.32	1.43	3.34	-	<del>-</del>	-	+
744933	AA625812	2.15	1.01	0.82	2.14	2.63	+	-	+	•
321386	1	2.45	1.56	0.77	1.57	3.19	-	-	-	+
253 <b>577</b>		1.48	1.24	0.41	1.19	3.57	-	•	-	+
41174		4.02	2.38	1.31	1.69	3.07	-	•	•	+
149013		7.82	5.75	2.30	1.36	3.40	-	-	+	+
296529		9.83	7.09	2.92	1.39	3.37	-	-	-	+
	AA017200	5.63	3.33	1.84	1.69	3.06	-	-	-	+
154749		1.76	1.07	0.57	1.65	3.10	-	-	-	+
197176		20.79	10.24	7.65	2.03	2.72	-	-	-	+
	AA 1903 13	24.22	21.29	6.72	1.14	3.61	-	<del>-</del>	+	+
	AA598561	14.33	5.93	6.17	2.42	2.32	-	+	-	•
	AA147837	10.43	6.02	3.47	1.73	3.00	-	-	-	+
	AA042928	9.14	5.47	2.99	1.67	3.06	-	•	-	+
	7 AA872122	85.21	65.01	24.92	1.31	3.42	•	•	+	-
28319	1	3.80	2.01	1.34	1.89	2.84	-	•	+	
	2 AA126708	18.65	13.11	5.65	1.42	3.30	-	•	+	+
	SAA873060	6.33	4.83	1.85	1.31	3.41	-	-	+	+
	1 AA479154	6.80	4.46	2.13	1.53	3.20	-	-	+	+
32560		7.30 14.02	3.27 6.63	2.93 5.40	2.23	2.49	-	-	-	+
3207	9 AA455917 6 R17320	1.02	0.03	0.54	2.11 2.79	1.92	•	+	_	T .
	0 AA777781	7.53	4.62	2.45	1.63	3.08	-	_	_	+
	0 AA962236	6.44	4.02	2.43	1.50	3.08 3.21	_	_	+	+
	7 AA664155	8.56	4.10	3.27	2.09	2.62	-	_		+
	2 AA677185	7.68	5.14	2.40	1.49	3.20		-	+	+
2661		× 1.94	0.59	1.40	3.31	1.39	-	+	-	-
	9 AA988701	24.27	24.53	6.55	0.99	3.70	-	-	-	+
34272	1	14.93	6.70	6.06	2.23	2.46	-	-	+	+
	45 AA 156202		5.11	1.14	0.85	3.84	_	-	+	+
	50 AA488481		1.90	2.57	2.69	1.99	-	+	•	-
2822			1.00	0.22	0.84	3.85	-	-	-	+
2031			11.94	3.48	1.06	3.63	-	-	+	+
403	38 R53072	2.61	1.86	0.79	1.40	3.28	-	-	-	+
2041	46 H55920	2.40	1.26	0.86	1.90	2.78	-	-	+	-
	83 AA431407		0.85	0.42	1.55	3.13	-	-	+	-
1419	72 R68909	18.88	10.40	6.60	1.82	2.86	-	-	-	+
3568			0.94	2.45	3.38	1.30	-	+	-	-
	:66 AA033564		1.68		1.62	3.05	-	-	•	+
2466			15.47		1.09	3.58	-	-	-	+
	21 AA48568					3.87	-	-	•	+
	32 AA28433					3.05	-	-	-	+
426	559 R6112	6 10.21	3.36	6.30	3.04	1.62	-	+	-	•

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					Table 1-3					
460841	AA708189	1.02	0.58	0.35	1.75	2.91	-	-	-	+
306771	N91887	3.61	1.62	1.49	2.23	2.42	-	-	-	+
160485	H22136	5.17	2.96	1.78	1.74	2.90	•	-	+.	+
	AA599107	38.86	34.73	11.02	1.12	3.53	-	-	+	+
75886	T59478	14.08	9.92	4.36	1.42	3.23	-	-	+	+
283919	N50797	7.56	4.94	2.43	1.53	3.11	•	-	+	+
813271	AA455938	0.96	1.12	0.25	0.85	3.79	•	-	•	+
	AA490561	3.25	1.81	1.14	1.79	2.85	•	-	•	+
242706	H93552	5.15	2.96	1.78	1.74	2.90	•	•	-	+
	AA775957	3.16	1.23	1.52	2.56	2.07	•	+	-	-
	AA446690	6. <b>96</b>	2.63	3.51	2.65	1.98	•	+	-	-
	AA042858	6.24	5.89	1.75	1.06	3.57	•	-	+	+
	AA629524	5.38	3.61	1.71	1.49	3.14	•	-	-	+
26406	R20638	6.56	5.37	1.92	1.22	3.41	•	-	•	+
	AA883160	4.29	2.67	1.42	1.61	3.02	•	-	-	+
	AI004930	39.75	3 <b>8.39</b>	11.08	1.04	3.59	-	-	-	+
	AA599120	9.36	3.14	5.71	2.98	1.64	-	+	-	-
	AA463200	9.64	6.52	3.08	1.48	3.13	-	-	-	+
	AA669183	4.20	2.32	1.50	1.81	2.80	•	-	+	+
	AA456271	11.89	6.59	4.25	1.80	2.80	-	-	-	+
	AA074224	2.79	1.08	1.39	2.58	2.02	-	+	-	-
	AA699770	3.42	4.00	0.91	0.85	3.74	-	-	+	+
324342		4.41	4.29	1.24	1.03	3.57	•	-	•	+
	AA029295	3.09	1.27	1.44	2.44	2.15	-	+	-	+
	AA489343	28.44	25.71	8.17	1.11	3.48	•	-	-	+
129725		17.94	10.53	6.22	1.70	2.88	•	-	-	+
121196		22.80	22.77	6.36	1.00	3.58	•	-	+	+
	AA457235	3.58	4.32	0.95	0.83	3.76	•	-	+	+
283142	t	3.58	1.94	1.31	1.84	2.74	-	-	-	+
1636707 123980	1	19.22 2.38	12.06 0.92	6.43 1.19	1.59 2.59	2.99 1.99	-	- +	-	+
253009	•	115.86	35. <b>75</b>	86.54	3.24	1.34	-+	+	-	•
	AA705058	0.92	0.61	0.30	1.51	3.07	-		-	- +
	AA630584	1.30	0.82	0.30	1.57	3.00	-	-	+	-
51151		3.26	3.41	0.43	0.96	3.61	_	_	+	+
11940	•	1.22	1.22	0.34	1.00	3.57	_	_		+
	AA455931	4.14	2.24	1.53	1.85	2.71	_	_	-	+
21120		4.71	2.93	1.60	1.61	2.95	_	-	•	+
	1 AA412053	34.32	29.87	10.07	1.15	3.41	1 .	_	_	+
	0 AA873577	35.19	14.23	16.93	2.47	2.08	١.	+	-	+
	8 AA 197344	7.37	3.62	2.93	2.03	2.52	+	-	+	-
28928		5.44	3.58	1.79	1.52	3.03		-	-	+
	5 AA457717		1.79	1.20	1.83	2.72		-	+	
24306		9.34	5.76	3.20	1.62	2.92		-	+	-
76992	6 AA430615	9.93	5.82	3.51	1.71	2.83		-	+	-
28981		7.47	5.73	2.31	1.30	3.23		-	-	+
19634		122.93	41.22	79.12	2.98	1.55		+	-	-
81408	0 AA465424	8.19	4.01	3.29	2.04	2.49	-	-	+	-
7451		5.23	3.37	1.75	1.55	2.98		•	•	+
	3 AA412643	3.95	2.39	1.37	1.65	2.88		-	-	+
3486		2.74	2.06	0.85	1.33	3.20		-	+	-
	74 AA608514	28.90	11.69	14.05	2.47	2.06		+	•	-
7780		15.55	9.75	5.31	1.59	2.93		-	-	+
149230	04 AA894577	4.23	2.66	1.44	1.59	2.93		-	•	+
	•						-			

					Table 1-3		_			
263070	N20052	1.17	0.90	0.37	1.31	3.21	-	-	-	+
951216 A		2.02		0.67	1.50	3.02	-	-	-	+
308726	N93236	14.73		7.73	2.61	1.91	-	+	•	-
40136	R53348	5.23		2.77	2.62	1.89	+	-	-	-
1461528 A		3.95	2.84	1.27	1.39	3.12	-	•	+	+
	A424344	2.93	1.10	1.58	2.66	1.85	-	+	-	-
	A010223	3.43	1.72	1.36	1. <b>9</b> 9	2.52	-	• .	-	+
	A669443		16.78	8.14	1.47	3.03	-	-	-	+
744944 A	A625890	1.73 ·	0.90	0.67	1.92	2.59	-	-	• 1	+
202904	H54020	12.77	10.17	3.93	1.26	3.25	-	•	•	+
257382	N30699	12.80	10.78	3.86	1.19	3.32	-	•	+	+
590390	AA157150	6.42	3.41	2.45	1.88	2.62	•	-	•	+
460798	AA708152	1.30	0.65	0.52	2.00	2.50	-	-	-	+
815164	AA481135	0.88	0.71	0.27	1.25	3.25	-	•	-	+
214884	H74119	3.55	2.09	1.27	1.70	2.79	-	-	-	+
782439	AA431836	35.89		12.92	1.71	2.78	•	•	-	+
277596	N57002	2.91	1.45	1.17	2.00	2.49	-	-	+	-
344432	W73474	15.48	11.50	4.93	1.35	3.14	-	-	-	+
796341	AA459750	7.16	4.60	2.44	1.56	2.93	-	-	-	+
	AA416759	4.84	1.70	2.96	2.85	1.64	-	+	-	<del>-</del>
788518	AA452566	13.97	9.65	4.61	1.45	3.03	-	-	•	+
201628	R98008	7.51	5.63	2.39	1.33	3.14	-	•	+	+
	AA872057	35.80	21.64	12.69	1.65	2.82	-	-	-	+
	AA598595	1.79	2.04	0.50	0.88	3.59	-	•	•	+
	AA497010	15.69	7.46	6.62	2.10	2.37	-	+	•	+
	AA099855	33.49	26.22	10.49	1.28	3.19	-	-	+	+
	AA453713	0.52	0.34	0.18	1.52	2.95	•	-	+	+
	AA430205	2.67	1.22	1.18	2.19	2.27	-	-	+	•
	AA886333	3.68	1.85	1.49	1.99	2.47	+	+	_	_
31225	1	4.76	1.64	3.09	2.90	2.91	-	-	-	+
35807		2.91	1.90	1.00 0.66	1.53 1.75	2.70	-	-	•	+
	AA599094	1.77	1.02 2.92	1.82	1.73	2.74	-	_	-	+
	AA465116	4.97 13.21	2.92 6.75	5.32	1.76	2.48	- -	-	_	+
	AA453926 AA452933	9.50	6.13 6.97	3.09	1.36	3.08	_	-	+	+
	0 AA486516	14.94	10.35	4.99	1.44	3.00	_	-		+
	1 AA456975	52.82	30.51	19.51	1.73	2.71		-	•	+
	3 AA670353	7.20	6.82	2.13	1.06	3.38	l -	_	+	+
	8 AA775223	3.29	2.14	1.14	1.54	2.90	1 -	-	+	-
22252		2.88	2.10	0.94	1.37	3.05	-	-	-	+
	7 AA486300	2.15	1.48	0.73	1.46	2.96	-	-	-	+
	0 AA496800	6.35	2.19	4.18	2.90	1.52	-	+	•	-
	1 AA400247	1.20	1.08	0.36	1.11	3.31	-	-	-	+
5233		2.91	1.24	1.41	2.35	2.07		+	-	+
	8 AA148034		6.94	4.63	1.77	2.65	-	-	-	+
84062	20 AA488036		8.57	4.92	1.61	2.81	-	-		+
	99 AA630374		13.26	6.24	1.41	3.00		-	-	+
382:			1.95	0.94	1.44	2.98		-	-	+
	69 AA156940	1.41	0.77	0.55	1.82	2.59		<u>-</u>	-	+
3436	09 W69460		0.58	0.94	2.73	1.68	1	+	•	-
	96 AA286914		1.54	0.78	1.48	2.93		-	+	+
14932	52 AA886870		11.50	2.79	0.86	3.54		-	+	+
2033			0.74	1.05	2.57			+	<del>-</del>	+
4517	07 AA707659	7.59	5.80	2.45	1.31	3.09	7 -	-	+	+

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122747   T75072   37.20   31.66   11.55   1.18   3.22						Table 1-3					
815072 AA465180	22747	T75072	37.20	31.66	11.55	1.18	3.22		-		+
251731   H96904	815072	AA465180						-	•	. +	•
815563   AA456833   4.86   3.88   1.55   1.25   3.13   -   +   +				0.84	0.24			•	<b>=</b>	<u>.</u>	+
\$98210			4.86	3.88	1.55	1.25	1	•	•	•	+
SP8210   A.598597   8.13   8.02   2.42   1.01   3.35   -     +     +		AA 190747	1.58	1.06	0.55	1.49	2.88	•	-	-	+
48367				5.87		0.99	3.38	•	•	•	+
28436 R40663 0.89 0.37 0.46 2.43 1.93 +		<b>A</b> A59 <b>85</b> 97		8.02	2.42	1.01	3.35	-	-	-	+
259953   N32594   1.46   1.97   0.40   0.74   3.63   -		H09959				2.62	1.74	+	-	-	•
107822						2.43	1.93	+	•	-	-
713285 AA283029 0.89 0.56 0.32 1.58 2.78 + 1   195903 R92227 6.95 5.56 0.32 1.58 2.78 + + 1   195903 R92227 6.95 5.56 0.32 1.52 3.11 + 1   1501998 AA887201 2.03 1.50 0.68 1.36 3.00 + 1   1501998 AA887201 2.03 1.50 0.68 1.36 3.00 + + 1   1501998 AA887201 2.51 1.48 0.94 1.69 2.66 + +   1838478 AA457517 2.51 1.48 0.94 1.69 2.66 + +   1635655 AI016688 5.13 4.14 1.65 1.24 3.11 + +   1635655 AI016688 5.13 4.14 1.65 1.24 3.11 + +   149485 H16600 0.54 0.37 0.19 1.46 2.89 +   149773 AA043347 29.51 20.98 10.06 1.41 2.93 +   149771 AA043347 29.51 20.98 10.06 1.41 2.93 +   149777 R78725 71.81 48.67 25.09 1.48 2.86 +   1494971 R78725 71.81 48.67 25.09 1.48 2.86 +   1494971 R78725 71.81 48.67 25.09 1.48 2.86 +   149515 H08417 3.88 2.73 1.33 1.42 2.91 +   149797 AA443152 0.44 0.47 0.13 0.93 3.40 +   141978 W86378 24.58 18.77 8.16 1.31 3.01 +   1319113 R63022 4.32 3.44 1.41 1.26 3.06 +   1319218 R3345 AA47021 5.67 3.30 2.18 1.72 2.60 +   1319218 R3345 AA47021 5.67 3.30 2.18 1.72 2.60 +   1319218 R3345 AA47021 5.67 3.30 2.18 1.72 2.60 +   1319219 R3345 5.41 1.88 0.67 1.14 3.18 +   1319219 R3345 5.41 1.88 0.67 1.14 3.18 +   1319219 R3345 5.41 1.88 0.67 1.14 3.18 +   1319218 R3345 AA47021 5.67 7.30 2.18 1.72 2.60 +   1319219 R3345 5.41 1.88 0.67 1.14 3.18 +   1319219 R3345 5.41 1.88 0.67 1.14 3.18 +   1319219 R3345 5.41 1.88 0.67 1.14 3.18 +   1319218 R3345 AA47021 5.67 7.34 4.24 1.57 2.73 + +   1319218 R3345 AA47021 5.67 7.34 4.24 1.57 2.73 +   1319218 AA47021 5.67 7.34 4.24 1.57 2.73 +   141361 AA50190 AA47021 5.66 0.5							3.63	-	-	-	+
195903						1.69	2.67	-	-	-	+
42070   R60847   5.72   3.44   2.12   1.66   2.69   -							2.78	-	-	-	+
1501998								-	-	-	+
S03468								•	-	-	+
838478								-	-	-	+
281191								-	-	-	+
391984   A1003625								-	-	-	+
1635665       Al016688       5.13       4.14       1.65       1.24       3.11       -       -       +       +       +       429906       Ad033973       5.35       3.27       1.98       1.63       2.71       -       -       -       +       +       429906       Ad033973       5.35       3.27       1.98       1.63       2.71       -       -       -       +       +       429906       Ad033973       5.35       3.27       1.98       1.63       2.71       -       -       -       +       +       487773       AA043347       29.51       20.98       10.06       1.41       2.93       -       -       +       +       240961       H91011       2.11       0.82       1.20       2.57       1.76       -       +       +       240961       H91011       2.11       0.82       1.20       2.57       1.76       -       +       +       -       -       -       +       +       240961       H91011       2.11       0.82       1.20       2.57       1.76       -       +       +       -       -       -       +       +       420961       H404317       3.88       2.73       1.33       1.42								-	+	•	+
49485         H16600         0.54         0.37         0.19         1.46         2.89         -         -         +         +         429906         AA033973         5.35         3.27         1.98         1.63         2.71         -         -         +         +         487773         AA043347         29.51         20.98         10.06         1.41         2.93         -         -         +         +         701409         AA287948         1.54         1.24         0.50         1.23         3.10         -         -         +         +         240961         H91011         2.11         0.82         1.20         2.57         1.76         -         +         -         -         +         -         -         +         -         -         -         +         -         -         -         +         -         -         -         +         -         -         -         +         -         -         -         +         -         -         -         +         -         -         -         +         -         -         -         +         -         -         -         -         -         -         -         -         -		i						-	-	-	+
429906   AA033973   5.35   3.27   1.98   1.63   2.71   -							1	-	-	+	+
## ## ## ## ## ## ## ## ## ## ## ## ##								•	-	•	+
701409 AA287948								•	-	-	+
144977         R78725         71.81         48.67         25.09         1.48         2.86         -         -         -         +         +         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -								-	•	-	
240961 H91011 2.11 0.82 1.20 2.57 1.76 - + 1049033 AA778675 3.07 1.85 1.15 1.66 2.67 + + + 1784116 AA445743 0.26 0.10 0.15 2.61 1.72 - + + 1784116 AA445743 0.26 0.10 0.15 2.61 1.72 - + + 1784116 AA445743 0.26 0.10 0.15 2.61 1.72 - + + 1784116 AA445743 0.26 0.10 0.15 2.61 1.72 - + + 1784116 AA445743 0.26 0.10 0.15 2.61 1.72 - + + 1784116 AA445743 0.26 0.10 0.15 2.61 1.72 - + + 1784116 AA445743 0.26 0.10 0.15 0.93 3.40 + + 178416 AA445743 0.26 0.10 0.15 0.93 3.40 + + 178416 AA445743 0.26 0.10 1.88 1.63 1.31 3.01 + + 178416 AA129297 1.90 1.12 0.72 1.69 2.63 + + 178416 AA129297 1.90 1.12 0.72 1.69 2.63 + + 178416 AA15748 AA447021 3.63 6.81 2.82 1.50 - + + + 178416 AA447021 5.67 3.30 2.18 1.72 2.60 + + + 178416 AA447021 5.67 3.30 2.18 1.72 2.60 + + 178416 AA447021 5.67 3.30 2.18 1.72 2.60 + + 178416 AA447021 5.67 3.30 2.18 1.72 2.60 + + 178416 AA447021 5.67 3.30 2.18 1.72 2.60 + + 178416 AA447021 5.67 3.30 2.18 1.72 2.60 + + 178416 AA447021 5.67 3.30 2.18 1.72 2.60 + + 178416 AA447021 5.67 3.30 2.18 1.72 2.60 + + 178416 AA447021 5.67 3.30 2.18 1.72 2.60 + + 178416 AA447021 5.67 3.30 2.18 1.72 2.60 + + 178416 AA447021 5.67 3.30 2.18 1.72 2.60 + + 178416 AA447021 5.67 3.30 2.18 1.72 2.60 + + 178416 AA447021 5.67 3.30 2.18 1.72 2.60 + + 178416 AA447021 5.67 3.30 2.18 1.72 2.60 + + 178416 AA447021 5.67 3.30 2.18 1.72 2.60 + + 178416 AA447021 5.67 3.30 2.18 1.72 2.60 + + 178416 AA447021 5.67 3.30 2.18 1.72 2.60 + + 178416 AA447021 5.67 3.30 2.18 1.72 2.60 + + 178416 AA447021 5.67 3.30 2.18 1.72 2.60 + + 178416 AA447021 5.67 3.30 2.18 1.72 2.60 + + 178416 AA447021 5.67 3.30 2.18 1.75 1.43 2.88 + + 178416 AA447021 5.67 3.30 2.65 11.75 1.43 2.88 + + 178416 AA447021 5.67 9.66 3.34 1.11 3.19 + + 178416 AA700736 1.73 1.11 0.63 1.56 2.74 + + 178416 AA700736 1.73 1.11 0.63 1.56 2.74 -		l.						•	-	-	
1049033		)						•	-	-	+
45515 H08417 3.88 2.73 1.33 1.42 2.91 + + + 784116 AA446743 0.26 0.10 0.15 2.61 1.72 - +								-	+	-	-
784116								-	-		
796774 415978 W86378 24.58 18.77 8.16 1.31 3.01 + 415978 W86378 24.58 18.77 8.16 1.31 3.01 + 415978 R14801 4.83 2.97 1.80 1.63 2.69 + 564846 AA129297 1.90 1.12 0.72 1.69 2.63 + 503097 AA151486 7.27 4.27 2.78 1.70 2.61 + - 139113 R63022 4.32 3.44 1.41 1.26 3.06 + + - 131621 R23790 10.21 3.63 6.81 2.82 1.50 - +								-	-	+	+
415978 W86378 24.58 18.77 8.16 1.31 3.01 + + 68636 T49801 4.83 2.97 1.80 1.63 2.69 + + 554846 R14631 2.29 2.07 0.71 1.11 3.21 + + 564846 AA129297 1.90 1.12 0.72 1.69 2.63 + + 503097 AA151486 7.27 4.27 2.78 1.70 2.61 + + 131621 R23790 10.21 3.63 6.81 2.82 1.50 - + + + 131621 R23790 10.21 3.63 6.81 2.82 1.50 - + + 56386 R73545 2.14 1.88 0.67 1.14 3.18 + + 784264 AA447021 5.67 3.30 2.18 1.72 2.60 + + 156386 R73545 2.14 1.00 1.27 2.41 1.90 + - + 1310493 W31074 7.56 7.25 2.32 1.04 3.26 + + - 310493 W31074 7.56 7.25 2.32 1.04 3.26 + + - 282144 N51883 30.16 18.11 11.43 1.67 2.64 + + - 2435341 AA700736 1.73 1.11 0.63 1.56 2.74 + + 252904 H88485 5.40 2.26 2.83 2.39 1.91 - + + 433573 AA701655 10.67 9.66 3.34 1.11 3.19 + + 433573 AA701655 10.67 9.66 3.34 1.11 3.19 + + 252904 H88485 5.40 2.26 2.83 2.39 1.91 - + + 252904 AA59888 11.55 7.34 4.24 1.57 2.73 + + 27168 AA521292 6.51 2.85 3.24 2.29 2.01 + - + + 27168 AA521292 6.51 2.85 3.24 2.29 2.01 + + + 4316106 AA701981 4.14 2.42 1.60 1.71 2.58 + + + 436106 AA701981 4.14 2.42 1.60 1.71 2.58 + + + 434388 AA629554 4.89 3.97 1.60 1.23 3.06 + + + + 884388 AA629554 4.89 3.97 1.60 1.23 3.06 + + + 884388 AA629554 4.89 3.97 1.60 1.23 3.06 + + 884388 AA629554 4.89 3.97 1.60 1.23 3.06 + + 884388 AA629554 4.89 3.97 1.60 1.23 3.06 + + 884388 AA629554 4.89 3.97 1.60 1.23 3.06 + + 884388 AA629554 4.89 3.97 1.60 1.23 3.06 + + 884388 AA629554 4.89 3.97 1.60 1.23 3.06 + + 884388 AA629554 4.89 3.97 1.60 1.23 3.06 + + 884388 AA629554 4.89 3.97 1.60 1.23 3.06								-	+	•	-
68636         T49801         4.83         2.97         1.80         1.63         2.69         -         -         -         +         +         35812         R14631         2.29         2.07         0.71         1.11         3.21         -         -         -         +         +         564846         AAL29297         1.90         1.12         0.72         1.69         2.63         -         -         -         +         +         503097         AAL51486         7.27         4.27         2.78         1.70         2.61         -         -         +         -         -         +         -         139113         R63022         4.32         3.44         1.41         1.26         3.06         -         -         +         -         -         +         -         131621         R23790         10.21         3.63         6.81         2.82         1.50         -         +         -         -         +         +         -         -         +         -         -         +         +         -         -         +         +         -         -         -         +         -         -         -         -         -         +         -								-	-	-	
35812       R14631       2.29       2.07       0.71       1.11       3.21       -       -       +       +         564846       AA129297       1.90       1.12       0.72       1.69       2.63       -       -       +       +         503097       AA151486       7.27       4.27       2.78       1.70       2.61       -       -       +       -         139113       R63022       4.32       3.44       1.41       1.26       3.06       -       -       +       +       -         131621       R23790       10.21       3.63       6.81       2.82       1.50       -       +       -       -       +       +       -         156386       R73545       2.14       1.88       0.67       1.14       3.18       -       -       +       +       -         784264       AA447021       5.67       3.30       2.18       1.72       2.60       -       -       +       +         291706       N73480       2.41       1.00       1.27       2.41       1.90       +       -       +       -         784016       AA443698       33.80       23.65 <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td>-</td> <td>-</td> <td>-</td> <td></td>								-	-	-	
564846       AA129297       1.90       1.12       0.72       1.69       2.63       -       -       +       +         503097       AA151486       7.27       4.27       2.78       1.70       2.61       -       -       +       -         139113       R63022       4.32       3.44       1.41       1.26       3.06       -       -       +       +       -         131621       R23790       10.21       3.63       6.81       2.82       1.50       -       +       -       -       +       +       -         156386       R73545       2.14       1.88       0.67       1.14       3.18       -       -       -       +       +         784264       AA447021       5.67       3.30       2.18       1.72       2.60       -       -       -       +       +         291706       N73480       2.41       1.00       1.27       2.41       1.90       +       -       +       +         784016       2.44       7.56       7.25       2.32       1.04       3.26       -       -       +       +         784016       AA43698       33.80								•	-	-	
503097       AA151486       7.27       4.27       2.78       1.70       2.61       -       -       +       -         139113       R63022       4.32       3.44       1.41       1.26       3.06       -       -       +       +       +         131621       R23790       10.21       3.63       6.81       2.82       1.50       -       +       -       -       -       +       +       -       -       -       +       +       -       -       -       +       +       -       -       -       +       +       -       -       -       +       +       -       -       -       +       +       -       -       -       +       +       -       -       -       +       +       -       -       -       +       +       -       -       -       +       +       -       -       -       +       -       -       -       +       -       -       -       +       -       -       -       +       -       -       -       +       -       -       -       -       -       -       -       -       -       -								-	-	-	
139113							1	-	-	_	т
131621       R23790       10.21       3.63       6.81       2.82       1.50       -       +       -       -       +         156386       R73545       2.14       1.88       0.67       1.14       3.18       -       -       +       +         784264       AA447021       5.67       3.30       2.18       1.72       2.60       -       -       -       +       +         291706       N73480       2.41       1.00       1.27       2.41       1.90       +       -       -       +       -         310493       W31074       7.56       7.25       2.32       1.04       3.26       -       -       -       +       +         784016       AA443698       33.80       23.65       11.75       1.43       2.88       -       -       -       +       +         282144       N51883       30.16       18.11       11.43       1.67       2.64       -       -       +       -       -       +       -       -       +       -       -       +       -       -       -       +       -       -       -       -       +       -       - <t< td=""><td></td><td></td><td></td><td></td><td></td><td></td><td>1</td><td>_</td><td>-</td><td></td><td>-</td></t<>							1	_	-		-
156386       R73545       2.14       1.88       0.67       1.14       3.18       -       -       +         784264       AA447021       5.67       3.30       2.18       1.72       2.60       -       -       +       +         291706       N73480       2.41       1.00       1.27       2.41       1.90       +       -       +       -       -       +       -         310493       W31074       7.56       7.25       2.32       1.04       3.26       -       -       -       +       -         784016       AA443698       33.80       23.65       11.75       1.43       2.88       -       -       -       +       +         282144       N51883       30.16       18.11       11.43       1.67       2.64       -       -       +       -       -       +       -         282144       N51883       30.16       18.11       11.43       1.67       2.64       -       -       +       -       -       +       -       -       +       -       -       -       +       -       -       -       +       -       -       -       -		1						_	_	_	T .
784264		t .					1	_		_	+
291706         N73480         2.41         1.00         1.27         2.41         1.90         +         -         +         -         310493         W31074         7.56         7.25         2.32         1.04         3.26         -         -         -         +         +         -         +         +         -         -         +         +         -         -         +         +         -         -         +         +         -         -         -         +         +         -         -         -         -         +         +         -         -         -         +         +         -         -         -         +         +         -         -         +         +         -         -         +         +         -         -         -         +         +         -         -         -         +         +         -         -         -         +         +         -         -         -         -         +         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         <								_	_		
310493       W31074       7.56       7.25       2.32       1.04       3.26       -       -       +       +         784016       AA443698       33.80       23.65       11.75       1.43       2.88       -       -       -       +       +         282144       N51883       30.16       18.11       11.43       1.67       2.64       -       -       +       -       -       +       -         435341       AA700736       1.73       1.11       0.63       1.56       2.74       -       -       -       +       -       -       +       -       -       +       -       -       +       -       -       +       -       -       +       -       -       +       -       -       -       +       -       -       -       +       -       -       -       +       -       -       -       +       -       -       -       -       +       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -								+	-	+	•
784016 AA443698 33.80 23.65 11.75 1.43 2.88 + + 282144 N51883 30.16 18.11 11.43 1.67 2.64 + - + 435341 AA700736 1.73 1.11 0.63 1.56 2.74 + + 252904 H88485 5.40 2.26 2.83 2.39 1.91 - + + 433573 AA701655 10.67 9.66 3.34 1.11 3.19 + + + 43041 AA406040 6.48 3.98 2.43 1.63 2.67 + + 897971 AA598868 11.55 7.34 4.24 1.57 2.73 + 827168 AA521292 6.51 2.85 3.24 2.29 2.01 + - + - + - 141361 R64454 1.28 0.68 0.54 1.90 2.40 + - + - 767298 AA418383 19.37 16.76 6.18 1.16 3.14 + + - 436106 AA701981 4.14 2.42 1.60 1.71 2.58 + 884388 AA629554 4.89 3.97 1.60 1.23 3.06 +								-	-		+
282144 N51883 30.16 18.11 11.43 1.67 2.64 + - 435341 AA700736 1.73 1.11 0.63 1.56 2.74 + + 252904 H88485 5.40 2.26 2.83 2.39 1.91 - + + 433573 AA701655 10.67 9.66 3.34 1.11 3.19 + + + 43041 AA406040 6.48 3.98 2.43 1.63 2.67 + 897971 AA598868 11.55 7.34 4.24 1.57 2.73 + 827168 AA521292 6.51 2.85 3.24 2.29 2.01 + - + + - 141361 R64454 1.28 0.68 0.54 1.90 2.40 + - + - 767298 AA418383 19.37 16.76 6.18 1.16 3.14 + + - 436106 AA701981 4.14 2.42 1.60 1.71 2.58 + 884388 AA629554 4.89 3.97 1.60 1.23 3.06 +								_	-	-	+
435341       AA700736       1.73       1.11       0.63       1.56       2.74       -       -       +       +         252904       H88485       5.40       2.26       2.83       2.39       1.91       -       +       -       -         433573       AA701655       10.67       9.66       3.34       1.11       3.19       -       -       -       +         743041       AA406040       6.48       3.98       2.43       1.63       2.67       -       -       -       +         897971       AA598868       11.55       7.34       4.24       1.57       2.73       -       -       -       +         827168       AA521292       6.51       2.85       3.24       2.29       2.01       +       -       +       -         141361       R64454       1.28       0.68       0.54       1.90       2.40       -       -       +       -         767298       AA418383       19.37       16.76       6.18       1.16       3.14       -       -       -       +         884388       AA629554       4.89       3.97       1.60       1.23       3.06       - <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td>-</td> <td>-</td> <td>+</td> <td></td>								-	-	+	
252904       H88485       5.40       2.26       2.83       2.39       1.91       -       +       -       -       -       433573       AA701655       10.67       9.66       3.34       1.11       3.19       -       -       -       +       +       +       +       -       -       +       +       +       -       -       +       +       +       -       -       +       +       +       -       -       -       +       +       -       -       -       +       +       -       -       -       +       +       -       -       -       +       +       -       -       -       +       +       -       -       -       +       +       -       -       -       -       +       +       -       -       -       +       +       -       -       -       -       -       +       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -	43534	1 AA700736	1.73					-	-	-	+
433573       AA701655       10.67       9.66       3.34       1.11       3.19       -       -       -       +         743041       AA406040       6.48       3.98       2.43       1.63       2.67       -       -       -       +         897971       AA598868       11.55       7.34       4.24       1.57       2.73       -       -       -       +         827168       AA521292       6.51       2.85       3.24       2.29       2.01       +       -       +       -         141361       R64454       1.28       0.68       0.54       1.90       2.40       -       -       +       -         767298       AA418383       19.37       16.76       6.18       1.16       3.14       -       -       -       +         436106       AA701981       4.14       2.42       1.60       1.71       2.58       -       -       -       +         884388       AA629554       4.89       3.97       1.60       1.23       3.06       -       -       -       +			5.40	2.26	2.83			_	+	-	-
743041 AA406040 6.48 3.98 2.43 1.63 2.67 + 897971 AA598868 11.55 7.34 4.24 1.57 2.73 + 827168 AA521292 6.51 2.85 3.24 2.29 2.01 + - + - 141361 R64454 1.28 0.68 0.54 1.90 2.40 + - 767298 AA418383 19.37 16.76 6.18 1.16 3.14 + 436106 AA701981 4.14 2.42 1.60 1.71 2.58 + 884388 AA629554 4.89 3.97 1.60 1.23 3.06 +	43357	3 AA701655	10.67	9.66	3.34			-	-	-	+
827168     AA521292     6.51     2.85     3.24     2.29     2.01     +     -     +     -       141361     R64454     1.28     0.68     0.54     1.90     2.40     -     -     +     -       767298     AA418383     19.37     16.76     6.18     1.16     3.14     -     -     -     +       436106     AA701981     4.14     2.42     1.60     1.71     2.58     -     -     -     +       884388     AA629554     4.89     3.97     1.60     1.23     3.06     -     -     -     +			6.48	3.98	2.43		2.67	-	-	-	+
827168     AA521292     6.51     2.85     3.24     2.29     2.01     +     -     +     -       141361     R64454     1.28     0.68     0.54     1.90     2.40     -     -     +     -       767298     AA418383     19.37     16.76     6.18     1.16     3.14     -     -     -     +       436106     AA701981     4.14     2.42     1.60     1.71     2.58     -     -     -     +       884388     AA629554     4.89     3.97     1.60     1.23     3.06     -     -     -     +								-	-	-	+
141361     R64454     1.28     0.68     0.54     1.90     2.40     -     -     +     -       767298     AA418383     19.37     16.76     6.18     1.16     3.14     -     -     -     +       436106     AA701981     4.14     2.42     1.60     1.71     2.58     -     -     -     +       884388     AA629554     4.89     3.97     1.60     1.23     3.06     -     -     -     +								+	-	+	-
436106 AA701981 4.14 2.42 1.60 1.71 2.58 + 884388 AA629554 4.89 3.97 1.60 1.23 3.06 +						1.90	2.40	-	•	+	-
884388 AA629554 4.89 3.97 1.60 1.23 3.06 +					6.18	1.16	3.14		•	•	+
							2.58		•	-	+
46294 H09132 1.03 0.35 0.78 2.97 1.31 + + -		1					3.06	} -	•	-	+
	4629	4 H09132	1.03	0.35	0.78	2.97	1.31	+	+	-	•

					Table 1-3					
253253	H891 <b>04</b>	1.73	1.30	0.58	1.33	2.95	-	•	-	+
1031346	LA609106	3.21	2.11	1.16	1.52	2.76	-	-	-	+
429574	AA011445	3.77	1.23	3.11	3.06	1.21	-	+	-	-
52327	H23459	5.70	4.77	1.85	1.19	3.08	•	-	•	+
810741	AA457725	72.43	48.42	26.10	1.50	2.78	-	-	-	+
155838	R72215	2.53	1.96	0.85	1.29	2.97	-	-	+	-
135083	R33917	17.07	17.59	5.18	0.97	3.29	-	•	•	+
2714 <b>72</b>	N35020	11.80	7.84	4.28	1.50	2.76	-	-	-	+
29964	R14699	1.12	0.63	0.45	1.79	2.47	-	-	•	+
269300	N35863	10.64	7.57	3.73	1.41	2.85	-	-	-	+
	AA011593	3.12	0.86	4.85	3.61	0.64	+	+	-	-
	AA598795	4.06	2.82	1.45	1.44	2.81	-	<del>-</del>	-	+
	AA436031	11.89	8.38	4.20	1.42	2.83	-	-	-	+
260628	H97565	8.94	7.49	2.93	1.19	3.05	-	-	•	+
730412	AA469966	1.25	1.00	0.42	1.25	3.00	-	-	-	+
	AA491312	12.13	9.01	4.19	1.35	2.90	-	-	+	-
342211	W63789	1.92	1.55	0.64	1.24	3.00	•	-	-	+
	AA644345	15.29	12.54	5.07	1.22	3.02	-	-	-	+
	AA496252	2.29	1.38	0.89	1.67	2.57	-	-	+	-
	AA598987	41.12	27.32	15.07	1.50	2.73	-	-	-	+
141336	R63 <b>823</b>	2.12	1.03	0.97	2.05	2.18	+	-	+	-
263894	N28522	0.46	0.42	0.15	1.10	3.13	-	-	•	+
25 <b>2864</b>	H88424	1.20	1.00	0.40	1.20	3.03	-	•	-	+
291 <b>057</b>	N72115	1.34	0.61	0.66	2.18	2.04	•	+	-	•
	AA417805	6.05	3.90	2.27	1.55	2.67	-	-	-	+
132911	R26186	52.56	22.96	27.26	2.29	1.93	-	+	-	+
	AA099404	2.11	1.57	0.73	1.34	2.87	-	-	-	+
	AA678065	2.38	1.00	1.30	2.39	1.83	-	+	-	-
	AA947922	4.13	3.78	1.32	1.09	3.12	-	-	-	+
	AA464995	0.44	0.43	0.14	1.04	3.16	-	-	•	+
	AA495962	4.17	3.13	1.45	1.33	2.87	-	-	•	+
271076	1	4.31	3.13	1.53	1.38	2.82	-	-	•	+
	AA424045	4.05	3.18	1.39	1.27	2.91	-	-	+	•
	AA070997 AA457082	5.75	3.77	2.16	1.52	2.66	•	-	•	+
	AA161283	2.82 0.66	2.95	0.87 0.21	0.96	3.23	-	-	-	+
3638		3.19	0.64 3.07	1.02	1.04 1.04	3.14 3.14	-	•	•	+
	AA599099	9.75	5.67	3.98	1.72	2.45	<del>-</del>	•	+	T .
	7 AA598625	2.02	1.65	0.68	1.72	2.95		-		+
	7 AA463449	1.59	1.01	16.0	1.22	2.59	-	-	_	+
27204		6.87	3.74	2.95	1.84	2.33	_	_	_	+
32861		0.46	0.40	0.15	1.13	3.03	-	_	_	+
	8 AA705047	0.55	0.43	0.19	1.27	2.89	_	-	_	+
	2 AA989139	2.25	1.64	0.81	1.37	2.79	_	-	-	+
	2 AA287936	4.23	3.51	1.43	1.20	2.96	-	-	+	+
23433		3.16	2.04	1.21	1.55	2.61	_	-	•	+
25164		0.50	0.47	0.16	1.06	3.09	-	-	-	+
5089		7.95	7.40	2.58	1.08	3.08		-	•	+
	5 AA521108	2.24	1.62	0.81	1.39	2.77	-	-	-	+
3029		10.63	10.42	3.39	1.02	3.14		-	+	+
	8 AA872704	238.01	171.23	86.12	1.39	2.76		-	•	+
	AA669452	41.65	28.65	15.44	1.45	2.70	-	-	-	+
	79 A1016074	3.27	2.02	1.30	1.62	2.53	-	•	-	+
2021:		29.46	16.52	12.46	1.78	2.36	-	-	+	-
	•						•			

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					Table 1-3					
197821	R93744	5.95	4.32	2.15	1.38	2.77		-	+	+
	A620421	3.02	1.90	1.18	1.59	2.55	•	•	-	+
	A633577	1.40	0.50	1.04	2. <b>7</b> 9	1.35	•	+	• .	-
	<b>A</b> 487020	2.47	3.20	0.73	0.77	3.36	•	-	•	+
197765	R93767	8.83	3.78	4.92	2.34	1.79	+	•	•	•
125308	R05886	2.61	0.93	1.97	2.80.	1.33	-	+	-	-
323611	W44508	1.25	0.50	0.76	2.48	1.64	-	+	-	•
	AA668897	12.39	4.81	8.02	2.58	1.55	•	+	•	-
	AA629591	9.95	9.24	3.27	1.08	3.04	•	-	+	+
	AA453849	7.62	7.57	2.45	1.01	3.11	-	-	+	+
	AA485675	2.59	1.28	1.23	2.02	2.10	+	-	+	•
243321	H95088	2.79	1.90	1.05	1.47	2.65	•	-	+	-
	AA633658	2.75	2.77 1.94	0.88	0.99 1.13	3.12 2.98	•	•	+	+
26185	AA013099 R20616	2.20 1.15	0.80	0.7 <del>4</del> 0.43	1.13	2.69	<u>-</u>	•	•	+
	AA027912	4.95	4.33	1.67	1.14	2.97		-	•	+
	AA487382	5.94	4.80	2.07	1.24	2.88	•	-	_	+
	AA001745	42.11	30.60	15.40	1.38	2.73	_	•	_	+
	AA504492	1.18	0.90	0.42	1.32	2.79	_	-		+
	AA629719	11.68	10.42	3.91	1.12	2.99	_	_	_	+
1	AA456717	6.60	2.40	4.88	2.75	1.35	+	-	_	· -
	AA206267	8.48	7.77	2.82	1.09	3.01	•	_		+
1	AA232645	0.85	1.07	0.26	0.79	3.30		_	-	+
	AA426629	3.80	3.11	1.32	1.22	2.87	_	_		+
	AA826324	3.06	2.56	1.06	1.19	2.90	-	-	-	+
246541	N77514	10.77	8.71	3.78	1.24	2.85	•	-	-	+
	AA401521	1.83	1.55	0.63	1.18	2.90	•	-	-	+
	AA706795	5.25	5.71	1.66	0.92	3.16	-	-	+	+
	AA521232	6.63	4.13	2.68	1.60	2.48	_	-	-	+
	AA677457	2.10	1.12	0.95	1.87	2.21	-	-	-	+
40075	R52015	3.13	3.07	1.03	1.02	3.05	-	-	-	+
127928		3.67	2.74	1.35	1.34	2.73	• .	-	-	+
279785	N48356	0.91	0.66	0.34	1.38	2.68	-	-	+	-
	AA983467	49.35	42.75	16.96	1.15	2.91	-	-	-	+
	AA482150	6.96	6.37	2.35	1.09	2.97	-	•	-	+
	A1014387	10.17	8.91	3.48	1.14	2.92	-	-	+	-
	AA663960	6.73	7.60	2.12	0.89	3.17	-	-	-	+
	AA873762	12.48	10.32	4.39	1.21	2.84	•	-	•	+
	AJ791165	3.87	3.96	1.26	0.98	3.07	•	-	•	+
	AA598874	26.99	22.87	9.42	1.18	2.86	-	-	<del>-</del>	+
	AA490734	4.08	3.05	1.51	1.34	2.70	i -	-	+	-
	AA196287	56.19	60.68	18.11	0.93	3.10		-	•	+
32887		3.04	2.20	1.15	1.39	2.64		•	•	+
	AA700604	28.82	38.07	8.83	0.76	3.26	•	-	+	- +
	AA464623	2.18	1.75	0.79	1.25	2.77	} -	-	-	+
244154		4.94	3.91	1.79	1.26	2.76	-	• -	•	+
	7 N62924 7 W15263	13.10	9.87	4.88	1.33	2.69	1	_	+	
	2 AA733038	3.1 <b>8</b>	2.15 4.35	1.26	1.48 1.55	2.53 2.46	1 -	_	-	+
	0 AA669124	6.74	4.33 2.43	2.74 1.78	1.55 1.69	2.40		<u>-</u>	-	+
	6 AA856556	4.11 30.68	28.07	10.55	1.09	2.91	]	-	•	+
	1 AA872153	30.68 14.80	6.17	9.24	2.40	1.60		+	-	-
8076		26.16	26.33	9.24 8.71	0.99	3.00			_	+
_	2 AA489194	4.46	3.05	1.76	1.46	2.53	1	•	•	+
02501	-11111303134	7.70	5.05	1.70	1.40	2.55	1			

				٦	Table 1-3		_			
51981	H23422	19.62	13.04	7.89	1.50	2.49	•	•	-	+
1591264 A	•	3.18	2.86	1.11	1.11	2.88	•	-	•	+
898265 A	A 508065	20.16	18.09	7.03	1.11	2.87	-	-	•	+
511459 A		37.04	46.42	11.64	0.80	3.18	•	-	•	+
296429	W00890	5.99	4.85	2.18	1.23	2.75	-	•	•	+
	A410480	1.78	1.58	0.63	1.13	2.85	-	-	-	+
	A456298	6.11	6.08	2.05	1.00	2.97	•	-	+	+
	AA402915	1.74	1.20	0.69	1.45	2.52	-	•	-	+
	AA504772	6.00	6.39	1.98	0.94	3.03	•	-	+	+
	AA878561	236.26	205.84	83.73	1.15	2.82	•	-	-	+
	AA827405	2.62	1.78	1.05	1.47	2.49	-	-	-	+
	AA421455	0.65	0.64	0.22	1.02	2.93	-		•	+
80292	T64433	2.18	1.64	0.83	1.33	2.62	-	•		+
	AA663826	9.12	11.76	2.88	0.78	3.17	•	-		+
75059	T50370		10.94	11.17	2.00	1.95	-	+	•	+
303139			1.22	1.00	1.77	2.17	-	-	+	•
48033			0.50	0.17	1.02	2.92	-	-	•	+
282720			6.24	3.18	1.33	2.61	-	-	•	+
291974			40.53	22.63	1.41	2.53	-	•	-	+
307029	3		44.33	21.22	1.27	2.66	•	-	•	+
	AA165402		6.33	2.13	0.99	2.93	•	-	•	+
	AA87335		79.29	37.33	1.26	2.67	•	-	+	•
	AA77996		0.91	1.09	2.13	1.78	-	+	•	- +
270 <b>7</b> 2			0.27	0.10	1.07	2.85	-	-	-	т
282895				2.03	1.99	1.93	+	-	+	+
	6 AA67934			1.23	1.47	2.45	-	-	-	+
87837	3 AA67033			10.19	0.87	3.05	•	<del>-</del>	-	+
	1 AA03494				1.75	2.16	-	+	•	+
	8 AA70242				1.19	2.72	-	•	- ,	+
86627	6 AA6631	50 1.94			1.06	2.84	-	•	-	+
24680			7 7.15		0.93	2.96	-	-	_	+
	52 AA 1338	96 1.3			0.84	3.05	-	-	_	+
4231					1.24	2.64	-	-	-	+
	17 AA4791	00 6.4			1.13	2.75	_	-	_	+
2340	36 H689				1.24	2.64	1 -	_		+
8843	33 AA6295	528 4.1			1.45	2.43 2.72		_		+
7583	47 AA4030	72 2.7			1.16 1.21	2.72	1	_	-	+
7957	38 AA4602					2.58		-	-	+
726						3.12		_	+	-
2517	753 H96	908 2.				2.08		-	+	-
9506	507 AA608	548 19.						-	-	+
	859 AA504							-	+	-
	456 H25							-	-	+
	577 AA490							-	•	+
	489 AA96							-	•	+
195								-	, <b>+</b>	-
	597 AA67							-	-	+
	5340 AA04			.82 6.7 :.10 0.5				-	-	+
	2806 AA40			10 0.3 ).78 40.4				-	-	+
	0935 AA43			). <b>87</b> 0.3				-	-	+
				1.57 0.8				•	+	•
		-		7.59 5.0				•	-	+
	6459 AA52			0.02 24.			71 -	•	-	+
86	8308 AA6	94006 6	J.77 U				•			_
										Р

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					Table 1-3					
	AA621031	1.92	0.89	1.16	2.15	1.65	+	-	-	-
486175	AA043133	10.39	9.02	3.91	1.15	2.66	-	•	+	-
81484	T63342	10.87	5.15	6.40	2.11	1.70	•	+		-
	AA007509	3.17	1.48	1.90	2.14	1.67	-	+	-	-
	AA446839	3.19	2.89	1.18	1.10	2.70	•	•	•	+
784128	AA446759	1.83	1.82	0.65	1.00	2.80	-	-	+	-
253865	N22007	1.68	1.49	0.63	1.13	2.67	•	-	+	+
	AA776810	2.05	1.00	1.17	2.05	1.75	-	+	-	-
595604	AA 167269	9.41	8.19	3.56	1.15	2.65	•	•	-	+
784129	AA432085	6.53	5.53	2.50	1.18	2.61	-	•	+	-
251452	H97993	3.12	2.79	1.17	1.12	2.67	-	•	+	-
767346	AA418557	4.50	3.43	1.82	1.31	2.47	•	-	+	-
138865	R62742	5.42	4.35	2.14	1.25	2.53	•	•	=	+
	AA427401	7.24	6.59	2.70	1.10	2.68	-	•	+	-
43705	H05706	0.35	0.33	0.13	1.07	2.71	-	•	-	+
76005	T47291	2.35	1.80	0.95	1.31	2.46	-	-	-	+
156045	R72518	2.01	1.50	0.83	1.34	2.43	-	•	. <b>-</b>	+
784085	AA446639	3.93	3.17	1.56	1.24	2.52	-	-	-	+
418053	W90732	7.57	7.14	2.80	1.06	2.70	-	-	-	+
322192	W37779	7.97	9.95	2.69	0.80	2.96	-	-	+	+
309864	W23847	2.49	1.62	1.12	1.54	2.22	+	-	+	-
44975		8.91	10.72	3.05	0.83	2.92	•	-	-	+
	AA504834	0.76	0.67	0.29	1.14	2.62	•	-	• -	+
291394	W03029	6.46	4.22	2.92	1.53	2.22	•	-	+	•
	AA461311	0.78	0.80	0.28	0.98	2.77	-	-	-	+
	AA480820	2.46	1.75	1.05	1.40	2.35	-	-	-	+
	AA045792	3.14	2.24	1.34	1.40	2.34	-	•	-	+
	AA629808	156.92	121.14	64.25	1.30	2.44	•	•	-	+
41411	4	112.58	99.23	43.25	1.13	2.60	•	•	-	+
247546		11.16	8.98	4.48	1.24	2.49	•	-	+	-
	AA496173	3.75	3.51	1.41	1.07	2.67	-	-	-	+
37665	li .	1.01	0.72	0.43	1.40	2.33	-	•	-	+
	AA113347	4.17	1.82	2.90	2.29	1.44	-	+	-	-
	AA292995	4.55	3.15	1.99	1.45	2.28	-	-	-	+
	AA902249	1.23	1.56	0.42	0.79	2.94	-	-		+
268178		28.02	27.47	10.37	1.02	2.70	-	-	-	+
	2 AA496148	5.14	3.76	2.18	1.37	2.36	•	-	+	•
	8 AA857413	249.73	233.03	94.34	1.07	2.65	-	-	-	+
	0 AA487934	4.05	3.61	1.56	1.12	2.60	-	•	-	+
	9 AA620404	2.48	1.95	1.02	1.28	2.44	•	-	-	+
	6 AA481438	17.53	6.47	17.49	2.71	1.00	+	•	-	-
	6 AA634164	6.77	5.15	2.83	1.32	2.39	•	-	+	+
	1 AA633768	54.39	52.93	20.36	1.03	2.67	•	-	-	+
	3 AA974971	1.21	0.85	0.53	1.41	2.28	•	•	+	- +
20678		6.05	5.71	2.30	1.06	2.63	-	•	•	+
	1 AA446479	0.73	0.44	0.35	1.64	2.05	•	•	-	T .
34505			0.35	0.71	2.47	1.22	-	+	•	+
	25 AA496736	2.58	3.15	0.90	0.82	2.87	-	•	-	+
	4 AA046321	2.58	2.44	0.98	1.06	2.63	-	-	•	+
7660	6 AA778869	5.15	4.43	2.04	1.16	2.52	} ~	-	-	+
	T51043	2.48	2.10	0.99	1.18	2.50		•	•	+
	5 AA454862	2.57	2.48	0.97	1.04	2.64		-	-	+
	37 AA167728	2.80 0.80	2. <b>07</b> 0. <b>57</b>	1.21 0.35	1.35 1.41	2.3 <b>2</b> 2.2 <b>7</b>	]	-	•	+
0,21,	// AA10//28	0.80	0.57	0.33	1.41	2.21		-	-	

				т	able 1-3						
122183	T98628	7.61	10.97 .	2 <i>.</i> 56	0.69	2.98	-	_	-	-	+
284545	N64762	1.10			1.42	2.24	-		-	-	+ ,
	AA485024	1.44			1.29	2.38	-		-	-	+
	AA192765	3.73	2.69	1.64	1.39	2.28	-		•	•	+
243784	N33927	2.99	2.12	1.33	1.41	2.25	-		-	-	+
	AA457529	41.97		15.95	1.02	2.63	-		-	•	+
	AA704459	1.29	1.51	0.46	0.86	2.79	-		-	-	+
83011	T70522	32.97		12.18	0.93	2.71	-		-	•	+
	AA679352	10.73	11.79	3.94	0.91	2.73	•		-	-	+
277761	N49605	2.43	1.65	1.12	1.47	2.17	-		•	-	+
	AA099593	4.03	4.49	1.47	0.90	2.74	-		-	-	+
	AA463822	1.35	1.17	0.55	1.15	2.48	-		•	+	-
33045		0.63	0.43	0.29	1.47	2.15	-		• .	+	-
137984	f .	1.63	1.42	0.66	1.15	2.47	-		•	. +	+
	AA968896	3.08	2.54	1.28	1.21	2.40	-		- '	-	+
138861		1.72	1.35	0.74	1.27	2.33	-		•	•	*
289645		0.92	0.67	0.42	1.38	2.21	-		-	+	-
140792		1.66	0.85	1.02	1.96	1.63	+		-	+	_
130130	R20813	0.29	0.22	0.12	1.29	2.29	-		-	+	- +
50374	I AA131466	3.64	4.28	1.33	0.85	2.74	-		•	Ŧ	+
	4 AA521243	2.52	2.65	0.96	0.95	2.63	-		•	•	+
	8 AA487650	11.96	14.60	4.33	0.82	2.76	-		-	-	+
	0 AA599178	12.52	17.40	4.38	0.72	2.86	-		-	_	+
45046	4 AA682819	3.97	2.31	2.13	1.72	1.86			•	_	-
	1 AA400214	3.84	1.71	2.87	2.24	1.34	+		-		+
	4 AA676920	0.80	0.68	0.33	1.17	2.41	] -		_		+
	74 AA630620	0.25	0.18	0.11	1.36	2.21 1.65	+		_		_
14232		1.49	0.78	0.90	1.92	1.05	+		_		-
	12 AA872143	224.84	89.33	214.43	2.52 1.48	2.09	1 :		_	+	-
	55 AA420992	2.04	1.38	0.98 5.51	0.88	2.68	1		-	_	+
	99 AA677499	14.78	16.78	3.51	0.88	2.63			_	-	+
2442		9.25	9.9 <b>7</b> 53.66	20.71	0.99	2.57			-	-	+
	68 AA456931	53.19 6.47	4.45	3.08	1.45	2.10			•	-	+
	68 AA774230	0.69	0.68	0.27	1.01	2.52	1	_	-	-	+
2514	H96504 251 AA598675	2.02	3.02	0.71	0.67	2.86		-	-	-	+
	68 AA779221	4.95	6.83	1.77	0.72	2.80		-	-	+	•
	722 AA455482		51.18	19.96	0.99	2.53	L	-	•	-	+
278:			74.35	20.99	0.77	2.74	4	<b>-</b> '	-	-	+
	905 AA625788		2.08	0.89	1.05	2.4		-	-	-	+
	727 H97851			2.17	1.26	2.2		-	-	•	+
	892 AA66945			3.19	0.99	2.5		-	-		+
487	373 AA046489			21.22	0.93	2.5		-	•	-	+
	863 AA48640			4.49	1.03	2.4		-	-	-	+
	988 AA41894				1.20	2.3		-	-	•	+
	7448 AA48947			12.76	1.06			-	-	-	+
	8642 AA97654		3.33		0.93			-	-	•	т
	6620 H1003				2.61			-	+	-	+
	5336 AA63037	6 5.6			1.08		40	-	-	•	+
	3087 AA97072	20 2.4					73	-	-	-	+
	1880 AA45463	34 3.1					72	-	-	-	
	1615 AA4876						.10	+	-	+	-
	9895 H007:						.13	+	-	-	-
46	1699 AA6822	74 3.4	1.3	5 3.84	2.5	0	.90	T			
											_

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					Table 1-3		_			
	AA609456	6.42	4.46	3.21	1.44	2.00		•	•	+
	AA490456	0.78	0.80	0.32	0.98	2.46	-	-	+	-
	AA699926	10.36	11.94	4.03	0.87	2.57	-	-	+	-
84078	T70999	0.34	0.26	0.16	1.29	2.14	-	-	-	+
415415	W81118	9.68	10.64	3.83	0.91	2.52	-	-	_	+
285226	N66278	25.74	40.08	9.23	0.64	2.79	-	•	+	-
41207	R56771	2.05	2.42	0.80	0.85	2.57	-	-	-	+
122239	T98662	6.50	5.28	2.98	1.23	2.18	_	•	-	+
284865	N66734	2.87	2.71	1.22	1.06	2.34	-	-	-	+
867717	AA780841	5.61	3.47	3.15	1.61	1.78	-	_	-	+
50562	H16903	34.20	30.09	15.14	1.14	2.26	-	-	+	-
71672	T58002	4.77	4.16	2.13	1.15	2.23	-	-	-	+
	AA135870	31.62	13.68	30.14	2.31	1.05	-	+	-	-
	AA017544	4.53	6.49	1.70	0.70	2.66	•	-	+	-
111389		2.59	2.15	1.21	1.20	2.14	-	•	-	+
	AA416782	0.75	0.75	0.32	1.01	2.32	•	•	-	+
	AA677574	1.64	1.62	0.71	1.01	2.30	-	-	+	-
488301	AA085748	0.97	1.24	0.38	0.78	2.52	-	-	+	-
	AA088371	7.14	7.60	3.02	0.94	2.36	-	-	-	+
	AA188999	5.86	3.92	3.24	1.49	1.81	-	-	-	+
	AA 598652	10.34	13.66	4.08	0.76	2.54	-	•	-	+
	AA457092	9.85	10.74	4.18	0.92	2.36	-	-	-	+
22600	1	0.36	0.18	0.28	1.99	1.29	+	-	-	-
343688		6.94	5.30	3 <b>.54</b>	1.31	1.96	-	-	+	-
	AA682399	8.65	14.46	3.24	0.60	2.67	•	-	-	+
	AA447503	3.99	2.02	3.12	1.98	1.28	-	+	-	-
845780	1	5.95	4.75	2.97	1.25	2.00	-	-	-	+
266500		7.14	7.39	3.12	0.97	2.29	-	-	-	+
119851		6.52	3.28	5.16	1.98	1.26	-	+	-	-
	AA455519	2.66	3.75	1.05	0.71	2.54	-	-	•	+
	AA432030	5.78	4.46	2.98	1.29	1.94	-	-	+	-
	AA206914	15.56	14.08	7.32	1.11	2.13	-	-	-	+
	AA279172	5.70	2.95	4.40	1.93	1.30	-	+	-	-
	AA782292	2.75	2.53	1.30	1.08	2.12	-	-	+	•
	AA 190997	11.04	13.42	4.64	0.82	2.38	-	-	+	-
	AA421769	15.96	26.86	6.13	0.59	2.61	-	•	+	-
	AA443966 AA477283	6.08	6.70	2.66	0.91	2.29	-	-	+	-
	2 AA001614	4.71	4.39	2.24	1.07	2.10	•	-	+	-
	AI000294	9.94	12.03	4.27	0.83	2.33	-	-	+	-
	3 AA001444	3.48 2.86	3.17	1.70	1.10	2.05	-	•	-	+
	8 AA486370	7.45	1.43	2.52	2.01	1.14	+	-	-	<del>-</del>
22006		4.81	7.83 8.77	3.42 1.92	0.95 0.55	2.18	-	-	-	+
	7 AA988630	13.38	13.90	6.58	0.96	2. <b>50</b> 2. <b>03</b>	-	-	- +	+
4128		0.72	0.52	0.45	1.38	1.59	_	-	+	-
	7 AA677706	24.08	105.26	8.92	0.23	2.70	<b>\</b>	_	+	+
15147		2.44	2.74	1.21	0.89	2.70	[	_	+	· ·
	0 AA001434	3.81	1.72	5.56	2.21	0.69		+	•	_
28983		6.67	4.11	5.46	1.62	1.22	+	_	•	•
	4 AA418670	8.86	14.76	4.05	0.60	2.19		•	+	_
12242		2.03	1.78	1.29	1.14	1.57	[	-	+	-
	4 AA406115	1.65	0.94	1.78	1.75	0.93	+	-		•
56344	4 AA112660	6.23	10.09	3.10	0.62	2.01		•	-	+
74094	1 AA478298	12.33	29.84	6.23	0.41	1.98		-	+	•
	1				3		1			

				Table 1-3	3				
462953 AA682423	3.21	5.03	1.90	0.64	1.69	-	-	+	-
1492202 AA875913	2.60	1.98		1.31	0.96		-	-	-

Table 1-4

					1 80(6							
<b>~</b>	GenBank			Aver.	Tumor	Tumor		771	T/N	T/B	T/B	T/B
Clone			Normals		Normal	/BPH	10X,>1/5	5X, >3/5	3X,>5/5	10X, >1/5	5X, >3/5	3X.>5/5
788180	AA453562	25.18	0.81	0.50	31.24	50.69	+	+	+	+	+	+
	AA196979	24.42	1.02	0.75	23.91	32.70	+	+	+	+	+	+
	AA779728	33.61	1.51	1.35	22 <i>.</i> 27	24.97	+	+		+		•
281003	N50880	128.77	6.30	4.90	20.43	26.26	+	+	+	+	+	+
308682	W25202	8.99	0.87	0.27	10.37	32.91	+	+	+	+	+	+
281039	N47717	14.05	0.94	0.90	14.99	15.57	+			+	·	•
284701	N64840	33.05	7.05	1.52	4.68	21.68				+	+	+
	AA490213	22.39	1.32	2.83	16.99	7.91	+	+	+	+	•	1
42415	R60981	6.86	0.51	0.72	13.52	9.49	+	+		+		
46166		13.13	0.93	1.71	14.12	7.68	+	+		+		
	AA432103	53.04	7.37	3.83	7.20	13.84		+		+		+
	AA427899	12.32	0.65	5.99	18.85	2.05	+	+	+	•		•
222157		8.80	1.02	0.72	8.67	12.14	+	+	+	+		,
48603 <i>5</i>	AA043109	24.75	3.47	1.86	7.13	13.32	+	+	+	+		+
450049	AA703396	18.37	1.83	1.80	10.05	10.21	+	+	+	+	+	+
66336	T66832	1.67	0.63	0.10	2.63	16.70		•	т .		+	+
35058	R45192	15.59	1.89	1.44	8.24	10.81	+	+	_	+		
26736	R39891	9.52	3.36	0.62	2.83	15.37	·	ν.	+	+		+
593185	AA159714	28.06	5.61	2.29	5.00	12.23	<b> </b>			+		+
32160	R43432	26.80	3.59	2.77	7.47	9.69		_		+		+
814053	AA465495	14.66	1.70	1.75	8.61	8.38	+	+	+	+	+	+
292515		16.22	2.86	1.56	5.67	10.38	ļ <sup>-</sup>	T		+		
<b>730</b> 739	AA436097	36.49	7.10	3.45	5.14	10.57	+		+	+	+	+
282089		11.18	1.51	1.41	7.41	7.92	l .		+	+	+	+
283312		63.94	6.63	12.14	9.65	5.27	+ +			+		
291255		15.20	1.35	4.65	11.29	3.27	ı	+	+			
	AA621342	20.15	3.04	2.59	6.63	7.78	†	+	+			
197525		9.52	1.85	1.06	5.16		+	+	+	+		+
67401		1.47	0.20	0.23	7.28	9.01				+		+
490178	AA121266	24.55	5.63	2.73	4.36	6.27	+			+		
	AA935560	0.88	0.15	0.12	5.96	8.99	1 +			+		+
	AA419486	9.35	2.23	1.07	4.18	7.33	+			+		
	AA780190	22.40	3.90	3.28	5.74	8.70	•			+		
259374		10.51	2.33	1.31	4.50	6.82		+	+		+	+
	AA437224	365.78	59.86	58.35		8.04				+		
	AA479284	23.83	3.20	4.92	6.11	6.27	+		+	+		+
	AA625634		4.34	3.34	7.46	4.85	+	+	+			
	AA282272		3.21	1.34	5.32	6.90		+	+			+
50772		18.10	6.74		3.59	8.58	† †			+		
	Al000271	6.58	4.38	1.91 0.63	2.68	9.47	1			+		+
	AA679414	10.34	1.11	4.04	1.50	10.44	<b>!</b> .			+	+	+
	AA047260		2.13		9.35	2.56		+	+			
1412481	AA845156	8.41	3.50	0.96	3.64	8.08	1			+		
283173				0.90	2.40	9.31	+			+		
	AA454963		3.97	2.47	4.49	7.21	4			+		
26519			2.97	7.43	8.28	3.30		+	+			
	AA235224		19.25	3.76	1.87	9.59				+	+	+
	AA232249		3.12	1.63	3.90	7.48	,				+	+
	) AA884926		1.54	0.16	1.07	10.14				+	+	+
258300			1.13	0.65	4.03	6.96	I.			+		
	AA682905		4.53	2.65	4.01	6.86	4			+		
	7 AA682905 3 AA991856		6.63	22.20	8.20	2.45		+	+			
1010440	סכפוגנעט י	135.55	57.06	16.44	2.38	8.25	1			+		-

				Table 1	4						
461522 AA705237	1.21	<b>u.65</b>	0.14	1.87	8.74				+		
868282 AA633957	8.99	2.01	1.47	4.48	6.12			+		+	+
	20.52	5.86	2.89	3.50	7.09	+			.+		
240694 H78135	4.53	0.90	0.83	5.01	5.47			+			+
321706 W35398	7.80	1.86	1.25	4.20	6.23				+		
770066 AA430545	11.43	2.60	1.93	4.40	5.94	+			+		
811603 AI732784	19.62	2.74	6.22	7.16	3.15		+	+			
51773 H23211	11.48	2.40	2.09	4.77	5.50			+			+
66317 T66815	5.50	1.76	0.78	3.13	7.06			_	+		
35626 R45292	9.42	1.09	6.51	8.67	1.45	+	+	+			
4503 <b>07</b> AA682851	9.39	1.62	2.25	5.80	4.17	+	+				
447715 AA702781	14.83	7.21	1.88	2.06	7.90				+		
731290 AA416843	11.60	4.56	1.57	2.54	7.40				+	+	+
825857 AA504783	4.27	2.51	0.52	1.70	8.20				_	7	•
565235 AA136125	35.83	12.59	5.15	2.85	6.96			_	+		
756600 AA481464	47.06	7.13	14.79	6.60	3.18	1	+	+			
293635 N63807	26.47	3.44	12.94	7.69	2.05	+	•	т.			+
757392 AA426110	0.86	0.37	0.12	2.32	7.41 8.16					+	· +
595695 AA173189	3.58	2.33	0.44	1.54 3.64	5.92				+	•	•
151055 H02230	27.32	7.51	4.62 3.27	6.12	3.41		+	+	•	-	
838366 AA457172	11.15	1.82	0.11	1.87	7.58		•	•	+	+	
460279 AA677629	0.87 3.07	0.46 2.45	0.11	1.25	8.17				+	+	+
595297 AA173598		0.93	1.77	6.13	3.21		+	+			
1631713 AI025015	5.68 12.35	3.96	1.99	3.12	6.21				+		
564801 AA129551 293651 N69653	13.35	1.84	6.56	7.26	2.04	+	+	+			
745347 AA625666	2.15	0.29	1.14	7.39	1.88		+	+			
772304 AA404486	113.14	18.10	38.24	6.25	2.96		+	+			
262053 N25030	10.75	5.15	1.52	2.09	7.08						+
81331 T60075	2.84	0.78	0.51	3.63	5.53	+			+		
840894 AA482340	34.81	11.87	5.66	2.93	6.15					+	
839081 AA488732	16.60	5.37	2.78	3.09	5.98				+		
796366 AA459761	34.94	17.03	5.05	2.05	6.92				+		+
359661 AA011095	26.24	8.79	4.39	2.99	5.97						+
50188 H17943	10.75	2.29	2.53	4.69	4.25			+			+
84955 T74846	5.26	1.01	1.43	5.21	3.69		+				
159608 H16152	14.43	4.14	2.69	3.49	5.36				+		
1390584 AA843592	6.54	1.91	1.21	3.42	5.41				+	_	
825736 AA504842	0.92	0.60	0.13	1.54	7.26					+	+
22161 T66154	20.24	13.40	2.80	1.51	7.24					+	
787938 AA452278	5.09	1.27	1.08	4.01	4.71		,	+		•	
809603 AA442991	49.03	6.94	29.52	7.06	1.66	+	+	т .	+		+
826109 AA521327		0.54	0.10	1.38	7.27	1			+		
884606 AA630006		13.42	5.43	2.49 1.48	6.16 7.16	İ			+		+
825822 AA491370		9.48	1.97	5.44	3.20	1	+	+			
531028 AA070495		3.89	6.62	1.72	6.83	1	•	·	+	+	
814062 AA465338		0.47 0.82	0.12 0.19	1.72	6. <b>93</b>	•				+	
826103 AA521416		1.72	2.71	5.19	3.28			+			
461670 AA779919		0.61	1.15	5.53	2.93		+				
415700 W85697 815163 AA481144		0.57	0.19	2.09		•				+	
810558 AA46456		3.00	4.01	4.83		1		+			
810220 AA46468	=	0.39		5.60			+				
359781 AA01093	=			3.13							
33701 AAV1093	_ ,.02					•					

					Table 1	1-4						
795265	AA451807	4.58	1.26	0.97	3.63	4.74				+		
	AA485333	2.49	1.26	0.39	1.97	6.38					+	
	AA488986	14.11	4.37	2.84	3.23	4.96						+
186918	H43317	3.04	1.10	0.56	2.76	5.38				+		
191904	H38880	7.67	1.44	2.75	5.32	2.79		+	+			
	AA476234	9.44	1.78	3.46	5.31	2.73			+			
	AA521394	1.20	0.83	0.18	1.44	6. <b>59</b>				+	+	
30428	R42061	4.29	0.99	1.16	4.35	3.68	+			+		
	AA620995	10.33	4.78	1.77	2.16	5.84				+		
251806	H9 <b>664</b> 7	2.94	1.83	0.46	1.60	6.39						+
305538	N89861	11.44	3.31	2.54	3.46	4.50					+	
52933	H29315	3.59	1.99	0.58	1.80	6.14						+
	AA490522	1.38	1.09	0.21	1.27	6.65					+	
	AA666366	85.06	37.74	15.20	2.25	5.60						+
	AA018675	16.82	3.11	6.91	5.41	2.43			+			
253132	H89036	1.40	1.21	0.21	1.16	6.66						+
233419	H77706	5.55	1.14	1.90	4.87	2.93		+				
	AA953229	5.30	1.51	1.24	3.52	4.28			+			+
	AA608636	3.00	1.87	0.48	1.60	6.18				+		+
	AA620591	6.84	6.68	1.01	1.03	6.75				+		
	AA427395	10.54	2.90	2.58	3.63	4.08						+
	AA486533	79.86	42.51	13.69	1.88	5.83				+		
268211	N36327	3.43	0.70	1.23	4.93	2.78	+					
	AA487914	15.85	7.18	2.88	2.21	5.50						+
	AA461092 AA458516	30.31	5.27	15.52	5.75	1.95		+	+			
	AA777488	10.13	5.34	1.75	1.90	5.78						+
150262	H01129	42.54	7.72	19.71	5.51	2.16		+	+			
	AA449837	13.66 7.66	3.31	3.87	4.12	3.53			+			
	AA045699	39.41	2.78 23.94	1.58	2.76	4.86						+
	AA136386	3.93	2.97	6.63 0.63	1.65	5.95						+
	AA054701	39.47	7.39	17.54	1.32 5.34	6.27 2.25				+		
	AA461467	6.21	4.35	1.01	1.43	6.16		+	+			
	AA465598	12.37	6.41	2.19	1.93	5.65				+		+
53022		1.34	0.70	0.24	1.90	5.63					+	
	AA055306	12.60	3.02	3.78	4.17	3.34	+					+
	AA481045	1.12	0.67	0.19	1.67	5.84	•					
51700		3.66	1.21	0.82	3.02	4.47					•	+
80050		8.15	3.84	1.52	2.12	5.35				+		,
594994	AA172372	14.61	10.98	2.38	1.33	6.14				•		+
795907	7 AA460346	5.39	1.25	1.71	4.31	3.15	+					,
824126	AA490611	0.59	0.37	0.10	1.58	5.87					+	
511952	2 AA100674	225.66	92.92	45.17	2.43	5.00				+		
	AA496957	9.40	3.54	1.97	2.65	4.76				+		
	AA432075	17,23	6.93	3.50	2.49	4.93						. +
	5 AA918370	401.67	285.67	66.91	1.41	6.00						+
	7 AA988746	9.00	4.61	1.65	1.95	5.45				+		
	7 AA996028	23.76	9.09	4.98	2.61	4.77					+	
	9 AA460675	26.07	23.22	4.18	1.12	6.24				+	+	
4930		4.88	0.86	2.90	5.68	1.68	+					
	8 AA436142	12.77	7.46	2.26	1.71	5.64	}				+	
20702		81.88	14.79	45.41	5.54	1.80	[	+	+			
	5 AA458747	1.34	1.36	0.21	0.99	6.35	]					+
163610	8 A1015679	3.58	1.45	0.74	2.47	4.85	ļ					+

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					lable 1	-4						
796754 A	A460719	0.66	0.83	0.10	0.80	6.50					+	+
771053 A		16.14	6.63	3.34	2.43	4.83						+
289071		37.48	15.32	7.78	2.45	4.82				.+.		
1637282	A1005515	4.01	2.58	0.70	1.55	5.71				+		
263697	H99672	4.23	0.93	1.58	4.57	2.68		+				
586780 A	A130677	1.92	1.75	0.31	1.10	6.12				+		+
416374	W86202	4.49	2.66	0.81	1.69	5.52				+		
375682 A	A032221	31.04	12.66	6.60	2.45	4.70						+
853968	A1791173	12.10	2.86	4.21	4.23	2.87		+				
287728	N75893	2.76	0.70	0.89	3.97	3.10	+					
842927	<b>NA48644</b> 5	4.24	3.59	0.72	1.18	5.88				+	+	
878281	<b>4A6702</b> 86	6.80	3.30	1.37	2.06	4.97					+	
359701 A	A011176	5.21	2.32	1.09	2.25	4.78						+
897807	AA598531	5.08	2.77	0.98	1.83	5.16					+	
	AA663983	50.50	12.03	18.24	4.20	2.77			+			
264895	N21170	7.86	1.63	3.70	4.82	2.12			+			
	AA504120	11.96	2.92	4.19	4.09	2.86			+			
	AA705041	0.78	0.61	0.14	1.28	5.64					+	
	AA777705	3.81	0.83	1.64	4.58	2.32	+					_
	AA422139	17.93	9.29	3.62	1.93	4.95					+	+
259973	N47829	1.66	1.95	0.28	0.85	5.99					•	+
290370	N62301	0.62	0.48	0.11	1.30	5.53						+
	AA425534	15.94	8.17	3.27	1.95	4.87 2.53	+					•
	AA873159	5.15	1.21	2.03 0.12	4.26 1.11	5.66	τ.					+
252314	H87175	0.69	0.62		0.61	6.16				+		
	AA677165	20.56	<b>33.95</b> 3.55	3.34 1.31	1.81	4.92				·		+
	AA976714	6.44 0.81	0.59	0.15	1.36	5.37						+
	AA150502	7.72	6.17	1.42	1.25	5.44				+		
139771	R62352 AA705376	8.32	6.33	1.55	1.31	5.38					+	
279482	N48804	8.66	3.25	2.16	2.67	4.00				+		
	AA490609	4.64	2.75	0.94	1.69	4.95					+	
	AA609323	38.01	9.06	15.59	4.20	2.44	+					
250869		123.92	32.54	44.10	3.81	2.81	+					
	AA779391	18.03	7.40	4.35	2.44	4.14					+	
	AA459100	18.28	10.68	3.77	1.71	4.85						+
77915		14.99	17.48	2.63	0.86	5.70				+		
260116		0.88	0.86	0.16	1.03	5. <b>52</b>						+
210405		34.02	7.84	15.49	4.34	2.20			+			
	AA278473	4.77	2.81	0.99	1.70	4.82	j					+
	7 AA491265	0.74	0.41	0.16	1.81	4.68	<b>\$</b>					+
813973	3 AA455689	3.24	0.88	1.18	3.70	2.75	1		+			
	7 AA504536	0.96	0.65	0.19	1.48	4.97	}					+
83880	7 AA457681	1.90	2.58	0.33	0.74	5.71	1					+
160776	5 AA989473	3.96	1.73	0.96	2.29	4.14	1					+
8616	0 T72336	2.10	0.74	0.59	2.82	3.58	1					τ-
145616	O AA862465	41.61	40.32	7.76	1.03	5.36	1			+		
89776	7 AA598470		0.49	0.98	4.25	2.12			+			
	1 AA883679		0.71	0.26	1.71	4.63	ŧ					++
28076	3 N50563		0.50		1.05	5.28				_		т
	13 AA430351		23.92		1.74					+		+
	24 AA171459		1.73		0.98							+
	19 AI004315		12.24		2.04		L L		+			'
20411	11 H55907	7 3.38	0.87	1.40	3.86	2.42	١ :		+			

				Table 1	1-4						
491184 AA137144	6.35	1.83	2.27	3.47	2.80	+					
824237 AA491249	74.89	31.45	19.54	2.38	3.83						+
24208 R38412	7.49	3.51	1.85	2.13	4.05						+
267419 N24973	0.97	0.96	0.19	1.01	5.16						+
731410 AA412250	0.53	0.58	0.10	0.92	5.24						+
869450 AA680244	51.58	35.87	10.98	1.44	4.70				+		
124252 R02439	6.06	4.74	1.25	1.28	4.85				+		
593658 AA160080	1.08	1.02	0.21	1.07	5.04						+
134969 R31789	6.81	4.59	1.48	1.48	4.58				+		
430465 AA680186	5.55	1.53	2.30	3.62	2.41	+					
796351 AA459765	2.34	1.80	0.50	1.30	4.72						+
549101 AA083485	12.81	2.43	17.02	5.26	0.75		+	+			
824376 AA489696	0.49	0.34	0.11	1.42	4.58					+	
453599 AA679569	4.23	2.23	1.03	1.89	4.10						+
5301 <b>85</b> AA083671	8.46	7.83	1.73	1.08	4.88					+	
773248 AA425773	21.90	19.94	4.51	1.10	4.86					+	
772416 AA405597	1.03	1.35	0.20	0.76	5.19						+
1456721 AA864875	12.82	5.49	3.58	2.33	3.58						+
50764 H17363	0.43	0.23	0.11	1.89	4.00						+
259462 N29545	3.69	1.65	1.02	2.24	3.62				+		
366815 AA029490	71.61	18.49	36.97	3.87	1.94	+					
731016 AA421256	0.85	1.00	0.17	0.85	4.96						+
197779 R93515	8.61	6.61	1.91	1.30	4.50				+		
1519515 AA902831	0.85	0.55	0.20	1.55	4.25						+
511428 AA126009	24.31	24.09	5.08	1.01	4.79						+
757206 AA443969	5.25	4.09	1.17	1.28	4.47						+
264465 N20522	0.77	0.84	0.16	0.91	4.79						+
609052 AA176607	0.87	0.87	0.19	0.99	4.63						+
1637343 AI015359	5.03	4.49	1.12	1.12	4.50						+
186768 H51914	9.43	5.75	2.39	1.64	3.95				+		
613237 AA181767	1.07	0.78	0.25	1.37	4.22						+
841302 AA487213	7.41	3.64	2.09	2.04	3.54				+		
251517 H96605	0.73	0.80	0.16	0.91	4.64						+
855624 AA664101	17.75	8.17	5.31	2.17	3.34				+		
841220 AA486723	1.46	1.23	0.34	1.18	4.32						+
1631194 AA994757	21.83	14.39	5.47	1.52	3.99					+	
251435 H97989	0.73	0.90	0.16	0.81	4.68						ŧ
767406 AA417911	0.44	0.45	0.10	0.98	4.45						+
745283 AA625567	7.94	5.76	1.97	1.38	4.02						+
815017 AA465258	14.81	9.71	3.84	1.52	3.85						+
346889 W79834	0.89	0.21	0.79	4.21	1.12		+				
823895 AA490490 1553306 AA934762	1.57	1.72	0.36	0.91	4.40						+
361922 AA001403	7.92	5.52	2.06	1.43	3.84						+
290378 N64508	9.18 17.10	5.49 17.02	2.55	1.67	3.59						+
1602619 AA988798	135.32	117.37	4.07 33.70	1.00	4.20						<del>-</del>
815800 AA485216	1.22	1.44		1.15	4.01						+
377672 AA056001	2.59	1.44	0.29 0.81	0.84	4.25				+	+	
256680 H96392	1.72	1.47	0.81	1.87	3.20				₹		_
823578 AA497118	0.56	0.53	0.44	1.17 1.06	3.88 3.98						+
878152 AA775431	4.20	3.49	1.10	1.06							+
685922 AA262559	3.59	3.70	0.90	0.97	3.82 3.99						+
786525 AA452113	1.46	0.42	1.00	0.97 3.46	1.46	_					•
1594019 AA988701	24.27	24.53	6.55	0.99	3.70	+					+
	,		5.55	4.77	5.70						

					Table	1-4		
282283	N52675	0.84	1.00	0.22	0.84	3.85		+
813271 A	A455938	0.96	1.12	0.25	0.85	3.79		+
461336 A	\A699 <b>770</b>	3.42	4.00	0.91	0.85	3.74		+
742581 A	A400247	1.20	1.08	0.36	1.11	3.31		+
161456	H25546	13.35	10.78	5.08	1.24	2.63	+	

Table 1-5

	CDI						
GI	GenBank	Aver.	Aver.	Aver.	Prostate	T/N	
Clone	Acc No.	of	of BPH	Po	Spec Exp	& T/BPH	Secreted
282089	N48259	1.51	1.41	11.18	+	+	•
646037	AA196979	1.02	0.75	24.42	+	+	+
745283	£A625567	5.76	1.97	7.94	+		-
30428	R42061	0.99	1.16	4.29		+	+
35058	R45192	1.89	1.44	15.59		+	+
46166	H09076	0.93	1.71	13.13		+	+
67401	T49326	0.20	0.23	1.47		+	+
240694	H78135	0.90	0.83	4.53		+	+
280217	N47941	5.86	2.89	20.52		+	+
292515	N68465	2.86	1.56	16.22		+	+
593185	AA159714	5.61	2.29	28.06		+	+
730739	AA436097	7.10	3.45	36.49		+	+
1592006	AA953229	1.51	1.24	5.30		+	+
32160	R43432	3.59	2.77	26.80		+	
42415	R60981	0.51	0.72	6.86		+	
50188	H17943	2.29	2.53	10.75		+	
51773	H23211	2.40	2.09	11.48		+	
81331	T60075	0.78	0.51	2.84		+	
222157	H85806	1.02	0.72	8.80		+	
281003	N50880	6.30	4.90	128.77		+	
281039	N47717	0.94	0.90	14.05		+	
283173	N45236	3.97	2.47	17.85		+	
308682	W25202	0.87	0.27	8.99		+	
450049	AA703396	1.83	1.80	18.37		+	
486035	AA043109	3.47	1.86	24.75		+	
488964	AA047260	2.13	0.96	7.77		+	
490178	AA121266	5.63	2.73	24.55		+	
712950	AA282272	3.21	1.34	11.53		+	
757435	AA437224	59.86	58.35	365.78		+	
784168	AA432103	7.37	3.83	53.04		+	
788180	AA453562	0.81	0.50	25.18	į	+	
814053		1.70	1.75	14.66	[	+	
823940		1.32	2.83	22.39	{	+	
868282	1	2.01	1.47	8.99	1	, +	
877835		4.34	3.34	23.06		+	
1034473	1	1.51	1.35	33.61		+	
1034738		3.90	3.28	22.40	Ì	+	
1048810		3.04	2.59	20.15	1	+	
1412481		3.50	0.90	8.41		+	
1556526		0.15	0.12	0.88	1	+	
26519		19.25	3.76	36.05	}	τ	+
35626		1.09	6.51	9.42	ł		+
51700		1.21	0.82	3.66	Ì		
52933		1.99	0.58	3.59	i		+
77915		17.48	2.63	14.99	1		+
80050		3.84	1.52	8.15	1		+
84955		1.01	1.43	5.26	1		+
124252		4.74	1.43		1		+
134969		4.74	1.48	6.06			+
13490)	1 101/09	4.39	1.48	6.81	1		+

Table 1-5

	GenBank	Aver.	Aver.	A1	Descripto	TO CALL	
Clone	Acc No.	of	of BPH	Aver.	Prostate	T/N	
150262	H01129			10	Spec Exp	& T/BPH	Secreted
151055		3.31	3.87	13.66			+
159608	H02230	7.51	4.62	27.32			+
186768	H16152	4.14	2.69	14.43			+
186918	H51914	5.75	2.39	9.43			+
233419	H43317	1.10	0.56	3.04			+
251806	H77706	1.14	1.90	5.55			+
252314	H96647	1.83	0.46	2.94			+
256680	H87175	0.62	0.12	0.69			+
	H96392	1.47	0:44	1.72			+ .
259462	N29545	1.65	1.02	3.69			+
260116	N32044	0.86	0.16	0.88			+
263697	H99672	0.93	1.58	4.23			+
280763	N50563	0.50	0.10	0.53			+
283312	N54763	6.63	12.14	63.94			+
284701	N64840	7.05	1.52	33.05			+
290370	N62301	0.48	0.11	0.62			+
290378	N64508	17.02	4.07	17.10			+
291255	N72215	1.35	4.65	15.20			+
293635	N63807	3.44	12.94	26.47			+
293651	N69653	1.84	6.56	13.35			+
321706	W35398	1.86	1.25	7.80			+
346889	W79834	0.21	0.79	0.89			+
361922	AA001403	5.49	2.55	9.18			+
375682	AA032221	12.66	6.60	31.04			+
377672	AA056001	1.39	0.81	2.59			+
416374	W86202	2.66	0.81	4.49			+
430465	AA680186	1.53	2.30	5.55			+
450307	AA682851	1.62	2.25	9.39			+
454317	AA677165	33.95	3.34	20.56			+
461336		4.00	0.91	3.42			, <del>+</del>
461476		0.61	0.14	0.78			+
491184		1.83	2.27	6.35			+
491764	•	0.59	0.15	0.81			+
509731		23.94	6.63	39.41			+
509887	AA054701	7.39	17.54	39.47	}		+
511428		24.09	5.08	24.31			+
511952 530185		92.92	45.17	225.66	1	•	+
564801	B .	7.83	1.73	8.46			+
586780		3.96	1.99	12.35			+
593658		1.75	0.31	1.92			+
594 <b>99</b> 4		1.02	0.21	1.08			+
595695		10.98	2.38	14.61	1		+
609052		2.33	0.44	3.58			+
610124	1	0.87 1.73	0.19	0.87	ł		+
666334			0.32	1.70	1		+
731410	1	1.54	0.16	1.64	1		+
756600		0.58	0.10	0.53	1		+
757392		7.13 0.37	14.79	47.06	1		+
131392	AA420110	0.5/	0.12	0.86	1		+

Table 1-5

1	GenBank	Aver.	Aver.	Aver.	Prostate	T/N	
Clone	Acc No.	of	of BPH	of	Spec Exp	& T/BPH	Secreted
767406	AA417911	0.45	0.10	0.44			+
768643	AA430351	23.92	9.10	41.57			+
770066	AA430545	2.60	1.93	11.43			+
773248	AA425773	19.94	4.51	21.90			+
773479	AA427899	0.65	5.99	12.32			+
784130	AA432075	6.93	3.50	17.23			+
787938	AA452278	1.27	1.08	5.09			+
788655	AA449837	2.78	1.58	7.66			+
795265	AA451807	1.26	0.97	4.58			+
796161	AA461092	5.27	15.52	30.31			+
796239	AA460675	23.22	4.18	26.07			+
8115 <b>8</b> 2	AA458516	5.34	1.75	10.13			+
811603	A1732784	2.74	6.22	19.62			+
811867	AA454963	2.97	7.43	24.57			+
815163	AA481144	0.57	0.19	1.20			+
823615	AA496957	3.54	1.97	9.40			+
823895	AA490490	1.72	0.36	1.57			+
824126	AA490611	0.37	0.10	0.59			+
825822	AA491370	9.48	1.97	14.07			+
826072	AA521394	0.83	0.18	1.20			+
826109	AA521327	0.54	0.10	0.75			+
838285	AA458747	1.36	0.21	1.34			+
838807	AA457681	2.58	0.33	1.90	ł		+
839081	AA488732	5.37	2.78	16.60	Ì		+
840894	AA482340	11.87	5.66	34.81	1		+
855624	AA664101	8.17	5.31	17.75			+
878152	P	3.49	1.10	4.20	1		+
884606	1	13.42	5.43	33.46	1		+
897767	1	0.49	0.98	2.07			+
951313	1	6.68	1.01	6.84	1		+
1031583		9.06		38.01			+
1056172 1456160	ľ	4.78		10.33	1		+
1472689	1	40.32 1.21	7.76 2.03	41.61 5.15			+
1535596	1	285.67		401.67			+
1610448	1	57.06		135.55	1		+
1631849		12.24		25.01	1		+
1031043	7 71004313	12.24	3.71	23.01	i		т

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

P 46 8 41 4 4 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	Falls of the No. 1804: Genes Upregulated 5 fold of nighter in at teast.		nimor samisle of 3 fold of higher in at least 2 tumor earlibres.
	VONT. PAR		. To element
	,	=	Ē

					Turner Bold (MPM125/A Fold (MPM123/A Fold (MPM123/A Fold (MPM126/A # of Tumor			A)1401011-7	EcidiMPM122/A	Fold(MPM123/A F	old(MPM126/A	# of Tumor
	IMAGE	Gen Bank	HODE	AvaTumor	Fold(AveTumor/Av	Fold(MPM120/A F	old(MPMPM)oio:	veBPH)	veBPH)	veBPH)	veBPH)	Samples
	Clone ID	Number			<b>⊕</b> BPH)			7	52.32	22.54	10.38	€
•	700480	44453340	1.96	133.47	67.88	149.63	165.70	2	£ 5	63.59	35.90	æ
- ‹	7001007	A 4 106079	2.38	114.00	47.98	65.24	24.62	11.47	5.28	41.24	18.50	•
٦,	284701	N64840	2.41	73.10	30.37	31.27	20.07	5 5	15.57	45.42	20.45	æ
, 4	450048	AA703396	2.55	78.64	30.08	38.52	11.57	60.00	19.72	21.41	63.61	•
r 10	245768	N55268	0.01	0.21	22.49	11.34	98.80 26.83	22.70	15.57	11.61	13.29	<b>6</b> 0
<b>,</b> «c	281003	N50880	24.05	423.65	17.62	22.19	\$6.03	\$ 0 80 80	10.57	5.71	7.21	ထ
· ~	138805	R35051	2.58	44.38	17.34	29.78	20.14	51.60	9.24	10.23	2.96	ထ
- 60	133130	R28397	1.48	25.48	17.23	30.48	4. C. A.	3.85	4.31	3.32	7.01	<b>6</b> 0
, a	838688	AA467235	3.4	46.10	13.38	11.55	30.20	11.99	10.01	17.09	10.66	æ
2	179753	H51549	0.01	0.12	11.81	8.28	1.04	11.83	6.14	4.27	15.03	ဇာ
: <b>=</b>	1030855	AA821761	1.24	12.57	10.16	7.89	15.01	8.58	7.12	8.55	19.24	<b>6</b>
12	61433	H20747	2.07	19.58	9.43	6.78	46.40	7.85	4.72	3.65	6.31	ဗ
5	306086	N91003	0.80	<b>3</b> .	8.70	13.54	5.5	8.91	5.51	12.71	6.41	<b>o</b>
7	195975	R91398	0.50	3.83	7.67	5.89	6.00 10.05	6.80	7.18	5.75	8.20	9
5	21831	T72543	0.29	1.97	6.87	3.24	5.5	7.14	10.04	4.07	5.11	80
9	292217	N79180	0.08	0.42	6.73	96. G	10.11	800	5.13	5.23	11.57	89
4	825859	AA504784	1.91	12.19	6.37	6.28	<b>5</b>	8.60	4.79	4.07	3.73	60
<u>~</u>	428816	AA004681	0.10	0.60	6.03	6.54	<b>†</b> 6	20.00	7.48	3.67	8.48	60
<b>6</b>	185034	R88764	0.07	0.14	5.84	3.19	0.01	88	8.86	3.53	3.06	ø
29	359701	AA011178	2.91	12.87	4.42	3.21	3.01	4.18	3.84	3.52	3.85	89
2	243641	N49548	1.55	8.3 <del>4</del>	4.09	3.17	3.80	100	4.47	3.77	4.01	60
ដ	262036	H98688	0.22	0.88	3.95	3.77	5 4	18 94	12.53	19.49	9.01	ĸo.
23	287528	N62096	0.30	3.88	12.94	15.20	26.16	14.64	17.70	1.2	3.82	vo.
*	784168	AA432103	15.97	194.72	12.19	79.6	27.30	10.24	5.27	13.28	1.75	vo
32	250374	N31952	1.29	13.52	10.47	7.91	64.38 0.06	4	3.08	33.98	88.	ĸ
28	210820	H67712	0.19	1.82	9.61	2.75	9	7.67	13.10	1.22	3.27	<b>1</b> 0
22	183923	R83407	17.59	158.73	8.02	10.87	10.00	3.78	3.02	4.81	0.54	ĸ
28	37880	R61372	0.09	0.74	6.39	9.46	10.08	6.72	12.64	2.83	7.06	iO.
28	1031599	AA609485	0.27	2.07	7.85	80.C	3.5	4.03	15.83	2.66	4.49	ro.
ဗ္ဗ	86160	172338	1.71	11.25	6.59	96.7	3.86	13.77	2.12	4.36	8.18	κo
દ	268211	N29986	0.57	3.71	6.48	S S S						

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	INAGE Clone ID	•	AveBPH	AveTumor	*oid(AveTumor/A eBPH)	Fold(AveTumoriav Fold(MPM120/A Fold(MPM121/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A # of Tumor ebpH) veBPH) veBPH) veBPH). veBPH) veBPH) veBPH) veBPH	Fold(MPM121/A veBPH)	Fold(NPM124/A	Fold(MPM122/A veBPH)	Fold(MPM123/A veBPH)	Fold(MPM125/) VeBPH)	A # of Tumor Samples
Ş	487787	Number	800	230	83	5.95	10.82	5.48	10.02	7.	3.81	-
; ;	90000	7 × 6 6 7 4 6 7	2 6	30.40	8.28	1.91	3.57	4.27	14.38	<b>3</b> .0	Į,	หา
3 3	811582	A4464607	9 6	21.22	5.92	9.12	4.48	4,71	8.04	2.69	8.45	ĸ
; ;	428436	AA004412	0.53	3.12	5.83	1.13	4.65	4.73	5.71	11.11	7.68	S
<b>2</b>	130138	R20813	8	0.20	5.62	7.23	8.80	5.30	5.38	96.9	2.08	ĸ
3 %	347588	WR1432	8	0.24	5.62	2.99	4.97	4.10	<b>1</b> .9	3.72	11.48	vo.
<b>9</b> 7	785729	AA460291	0.02	0,11	8.58	5.69	8.9	0.63	11.83	6.34	3.73	S
9	825785	AA505117	55	8.57	5.51	3.24	86.9	2.95	8.31	8.01	<b>4</b> .	40
\$	49838	H29290	0.33	9.	6.30	2.31	3.63	6.60	3.61	13.09	3.10	so
: =	417075	WR7823	0.83	85.4	5.30	2.79	3.52	3.68	4.21	8.09	œ 2.	vo
<b>; ;</b>	840751	AA486071	1.73	00.0	625	3.31	10.29	3.87	2.36	7.58	4.13	vo
. 3	241821	H93381	90.	5.43	5.10	3.95	6.61	5.92	8.06	2.79	3.31	ĸo
‡	29063	R40970	1.68	8.52	5.08	4.26	8.83	4.57	7.80	4.62	2.43	45
<b>.</b> 5	124128	R01348	1.25	6.16	4.92	2.53	60.9	3.89	4.46	8.48	5.11	<b>v</b> o
\$	238821	H65030	4.70	23.13	4.92	2.33	3.91	3.66	11.43	4.02	4.20	מי
**	134719	R28287	0.29	1.40	4.85	5.02	7.68	4.08	5.94	3.53	2.83	40
4	267745	N82244	3.81	18.41	4.83	2.38	4.28	3.75	20.0	3.82	5.12	٠c
3	271141	N34482	0.01	90.0	4.58	0.79	8.80	4.19	3.79	5.37	4.58	<b>V</b> O
S	1240414	AA788780	0.79	3.60	4.55	1.68	4.13	3.48	7.19	6.34	4.47	s,
<u>.</u>	307249	N93438	3.38	15.28	2.	9.25	5.31	0.40	5.38	3.28	3.63	'n
22	783849	AA443688	0.13	0.59	4.48	4.82	5.19	6.45	4.67	1.71	4.04	so.
2	897720	AA508082	0.32	1.38	4.37	3.68	3.50	2.84	3.83	6.46	5.83	50
3	283070	N51297	0.03	0.12	4.31	3.87	96.9	3.48	5.58	<u>z</u> .	4.46	<b>.</b>
8	285005	N21081	0.05	0.20	4.23	6.18	3.28	3.22	<b>6</b> .00	0.58	8.18	٠,
28	343987	W70234	8.51	35.37	4.16	6.16	7.98	3.20	3.85	0.24	3.50	<b>v</b> o
6	416866	W87281	0.46	1.88	4.03	2.15	7.06	3.45	3.09	5.14	3.31	ĸ
80	842548	AA486281	1.15	4.62	4.02	3.48	4.63	2.64	1.1	5.85	3.05	vo
8	40699	R55882	0.05	0.18	3.70	3.25	3.93	4.50	3.68	4.50	2.31	w ,
Ş	359661	A011096	23.32	84.82	3.64	4.70	6.56	3.16	3.28	3.48	<b>7</b>	ю
2	128230	R06307	0.03	0.10	3.50	3.19	3.36	3.88	4.68	4.99	0.89	ĸ
95	375662	AA032221	15.08	45.43	2.90	3.03	3.29	3.51	3.03	3.10	1.45	••
3	1468280	AA884928	0.65	22.38	34.18	18.89	1.69	59.34	105.24	0.80	18.00	•
Z	743828	AA634379	0.58	12.90	22.20	29.78	48.11	29.58	0.88	23.62	1.28	<b>+</b>
65	1350584	AA843592	7.	33.09	18.02	49.03	48.43	3.22	1.15	2.77	3.51	<b>~</b>
							,					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

		Gen Bank			Fold(AveTumor/Av Fold(MPM120/A Fold(MPM121/A Fold(MPM124/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A \$ of Tumor	V Fold(MPM120/A	Fold(MPM121/A	Fold(MPM124/A	old(MPM122/A F	old(MPM123/A F	old(MPM126/A veBPH)	Samples
	MAGE	Accession	AveBPH	AveTumor	•BPH)	veBPH)	veBPH)	V•BFH				•
	Clone	Number			•		0.81	4.	57.03	15.40	1.72	4
8	41647	R52794	0.40	8.34	15.84	CO:55	5	1 88	8.52	48.21	5.35	*
3 6	7777	AA625880	1.80	22.42	11.81	2.76	51.4	3 8	9 28	2.32	1.91	4
9 6	1024473	AA770778	0.13	1.36	10.50	22.49	23.74	3.60	13.07	1.00	1.00	4
3 8	487185	AA045074	0.01	0.0	8.98	13.49	10.45	5.65	F.34	4.84	2.78	•
8 6	4458474	AA867997	0.08	0.69	8.20	10.16	21.53	8 8	56.70	4.57	6.15	4
2 2	360156	AA013260	0.01	0.11	8.02	10.07	0.73	S. 2	10.46	1.64	2.63	*
3 :	144740	R76229	0.19	1.27	<b>6</b> .55	7.58	10.38	3 5	2.85	10.25	15.67	4
: 5	24915	R38066	0.93	5.97	6.42	3.07	2.31		08:80	2.35	3.20	4
2 7	593185	AA165678	9.70	60.24	6.21	9.37	96.21	5 OF	2.19	1.74	10.42	•
¥2	1056172	AA620995	11.61	70.00	6.03	99.9	90°C	K 43	5.02	1.51	2.67	<b>~</b>
2 %	782383	AA431407	0.95	6.65	5.84	8.95	77'LL		187	2.05	3.19	4
4 2	1031113	AA609914	0.01	0.07	5.65	16.58		1 10	8.43	5.07	2.77	•
: #	481541	AA705258	98.0	1.88	5.48	2.60	0.47	7 6	9 02	2.29	2.67	•
2 2	491708	AA115259	0.75	3.80	5.23	3.93	6.17	 	1.30	0.26	5.98	4
: 2	787838	AA452278	1.76	8.87	15.04	6.20	9.84	3 6	1 52	6.64	9.17	•
3 2	754034	AA479058	0.02	0.10	5.01	5.74	6.67	8 8	2 80	1.71	3.26	4
: 2	970590	AA683077	9.0	0.18	4.93	3.02	13.78	90.5	8 52	9.64	<b>6</b> .09	•
£	68787	T53389	80	90.0	4.91	1.66	3.59	10.83	11.89	6.25	2.97	*
3 2	292515	N68465	3.53	17.34	4.91	3.22	3.11	2.03	25.6	2.70	10.45	•
£ £	296210	N74392	4.0	2.17	4.80	3.19	3.94	0.30 6.30	7.12	1.57	11.65	*
8	40402	R52069	0.0	0.09	4.82	3.44	0.53	2 5	9 28	4.42	6.71	•
87	858153	AA633811	90.0	0.24	4.82	1.99	4.80		3.55	3.98	1.23	4
<b>.</b> 82	704687	AA282253	0.27	1.28	4.71	1.92	5/6 5/6	3 3	5.28	1.16	2.07	<b>~</b>
8	814308	AA459318	9.65	45.42	4.71	5.23	01.01	938	7.51	1.30	7,89	4
8	781233	AA446316	0.02	0.10	4.70	0.48	4.03	88.8	3.38	1.85	2.40	*
5	823794	AA490263	1.70	7. 2	4.68	3.60	(0.0)	33	1.01	1.63	4.68	*
8	260170	N32072	12.85	59.29	4.61	5.52	0.00	3.87	8.59	2.38	2.87	4
8	449309	AA777902	2.47	11.22	2.5	4.85	3.00	40.04 40.26	1.78	4.81	3.78	
\$	236155	H61758	0.98	4.35	4.51	<b>4</b> .	2.42	3.73	5.40	9.60	3.44	<b>~</b>
9	296679	W02258	<b>3</b> 5.0	4.15	4.41	CR.	(£.2)	65. 6	1.35	5.02	1.71	4
8	195387	R88915	0.47	2.07	4.38	5.34	0.40	7.7	1.35	3.96	5.65	•
25	428267	AA004828	1.41	6.20	4.38	2.17	24.0	4.23	10.05	2.98	1.97	4
83	391984	AI003625	4.42	19.15	4 2	3.25	4 37	3.15	9.71	1.81	1.53	*
8	51700	H22854	1.82	8.25	<b>4</b> 28		}					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	IMA (25		;	1	Fold(AveTumoriav Fold(MPM120/A Fold(MPM121/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A # of Tumor	Fold(MPM120/A	Fold(MPM121/A	Fold(MPH124/A	Fold(MPM122/A	Fold(MPM123/A F	old(NPM126/A	Samples
	Clone ID	Ē.	AveBPH	Avelumor	•BPH)	veBPH)	veBPH)	VeBPH)	(wage)			•
•		Jagun N	,	A2 40	71.7	5.48	6.00	2.50	3.26	4.67	2.90	•
8	177.	*****	9.6	200		1 07	3.35	4.59	7.43	8.18	1.26	*
5	210897	H68840	0.75	8 9	2 9	8.87	3.20	5.81	2.50	1.76	4.48	₹
102	808456	<b>VV44</b> 3080	8 1	0.70	0.4	 	90	3.63	4.72	4.28	2.07	<b>~</b>
103	361659	W96189	6.37	20.02	80°	3.68	2.48	<b>\$</b> .08	2.17	8.08	3.8	•
호	148225	H13668	1.72	5.5	70.4		321	2.71	10.52	0.69	3.20	•
50	417867	W90128	13.56	27.72	<b>5</b> 6	1.27	30.	3.96	5.13	8.63	2.31	*
8	436531	AA703046	<b>3</b>	E 6	₹ 6	. e	5.02	3.09	2.74	2.97	3.41	₹
107	1031348	AA809106	1.7	<b>3</b> 6	A 70 C	. 60	8 9	11.7	3.69	6.17	1.75	4
2	1567933	AA939275	r 6.0	8.7	- 00 c	80.0	76.	<u>.</u> 4	1.37	4.41	3.71	4
9	144747	R76247	S. 6.	) i	B 7	5.10	6.74	3.81	2.10	3.38	2.84	<b>~</b>
2	321386	W32272	1.23	£ ;	<b>1</b> 000	9 F	74.47	5.10	3.29	7.01	0.18	4
Ξ	725285	AA291742	0.03	0.10	3.78	<b>2</b> .3 <i>t</i>	3.53	Q. E	6.85	1.28	0.83	4
112	258404	N25883	0.28	1.07	37.6	. e.	6. 4	2.58	3.38	4.60	4.50	4
113	289892	N25920		8.68	3.70	C. 83	3 1.0	200	2.44	4.10	5.85	<b>4</b>
=	283885	N53378	0.63	3.42	3.80	\$ E	3	255	4.74	3.85	0.48	7
115	359781	AA011320	3.58	13.00	365	4.67	0.0 1 4 4 4	S 6	5.17	4.30	1.00	•
118	854701	AA630104	0.98	3.58	3.65	26.7	2 6	3.62	905	3.32	2.93	*
117	248886	HB2081	0.83	3.8	3.65	<b>.</b> 5	R !!	1 8	38 8	2.38	1.1	4
118	771053	AA427528	10.52	38.08	3.62	3.55	5.47	10.0	5.5	3.29	5.98	4
119	488010	AA045709	0.88	3:5	3.60	1.73	67.0 78.7	- <del>-</del> -	9 4	4.57	1.28	<b>~</b>
120	112482	T90980	11.33	40.58	3.58	1.21	18.4	100	3 6	2.78	4.53	*
121	80036	T82842	0.03	0.10	3.58	5.81	3.02	5 G	<b>7</b> .19	A 82	60	~
122	855832	AA664104	0.80	3.38	3.52	2.09	3.77	<b>9</b>	78.1	<b>4</b> 6	1.82	•
123	811914	AA54968	2.86	10.39	3.51	3.92	5.10	3.87	9.75 1.75	£:7 7	10. 4.4	
124	281371	N47886	0.38	1.25	3.49	5.39	3.0 <b>4</b>	8. 8.	71.7	75.7		
125	363823	AA704650	9.01	31.38	3.48	2.44	3.12	<b>3</b> 8.	5.1	3.48	870	• •
12	395	AA757401	2.20	7.68	3.48	2.26	5.31	3.00	¥.08	4.15	2.00	•
154	138607	PK1753	7	88.	3.47	1.74	4.81	3.80	3.23	4.18	2.85	<b>→</b> ·
1	289071	NATAOB	19.65	88	3.46	3.37	5.09	1.38	1.59	6.09	3.28	→ ·
5 5	480203	AA677461	0.55	1.89	3.45	1.61	4.23	4.13	4.28	4.93	1.55	<b>.</b>
	242081	103334	274	9.21	3.40	5.12	6.73	3.43	3.25	0.82	0.85	4
3 5	207008	HARSON	91 0	0.53	3.38	3.02	4.77	3.04	2.28	4.65	2.52	₹
÷	306830	AA768154	990	2.24	3.36	1.87	5.62	3.24	4.80	1.60	3.25	₩
	209381	H64085	78.0	3.25	3.35	2.04	3.28	4.07	1.86	4.90	3.98	<b>~</b>
}							•					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

		700				Ending Fold (MPM121/A Fold (MPM124/A Fold MPM122/A Fold (MPM123/A Fold (MPM125/A # of Tumor	Fold/MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A	Fold(MPM126/A	# of Tumor
	IMAGE	Accession	AveBPH	AveTumor	Fold(Ave i umorize	VeBPH)	veBPH)	v•BPH)	veBPH)	veBPH)	VeBPH)	and mrs
	Clone	Number					,	A 86	3.28	1.47	3.16	•
13	23897	R38364	0.38	1.24	3.28	1.98	J.81	9	3.37	3.97	2.04	4
135	1487905	AA883504	1.18	3.84	3.27	1.87	3.82	n d	660	4.48	1.12	*
138	1556526	AA835560	1.17	3.77	3.21	3.58	5.88	3.40	38	3.33	3.88	4
137	491418	AA115537	0.5	1.72	3.20	1.82	3.32	0 7 6	4 48	5.11	1.80	•
138	744001	AA629039	3.87	12.24	3.17	1.60	3.05	3.27	e e	3.55	1.84	•
130	293110	N63848	3.38	10,70	3.17	2.38	3.94	<del>1</del>	380	3.22	3.21	*
9	782047	AA429802	1.17	3.67	3.14	3.26	2.51	3.73	) i	3.10	4.59	4
3	454880	9 4 8 7 7 2 0 6	1.02	3.16	3.08	2.08	3.15	1.69	3.07	2 9	1.87	*
: \$	384293	AA709370	0.50	1.53	3.05	4.26	3.84	4.05	3.20	3.71	29	*
<u> </u>	7007	D44132	0.53	1.63	3.6	2.04	3.94	3.79	3.14	- a	85.4	4
₹ ;	and a	101 E F F F F F F F F F F F F F F F F F F		800	3.02	3.33	3.10	2.05	4.16	8 6	970	•
*	70554/	AA400500	3 7	3.44	3.02	2.11	2.81	3.08	3.17	3.53	9 70 6	· ¬
2	09/971	K10013		3.83	2.98	3.22	3.29	1.89	3.35	7.77		• •
<del>2</del>	428288	AA007370	07:1	2 .	7	3.15	2.95	3.25	3.12	1.85	<b>1</b> .00	, ,
147	136819	R36465	0.47	1,3/	, t.	3.88	6.22	3.12	1.2	0.47	3.55	₩ .
<b>₹</b>	433350	AA700604	42.83	122.18	7.50	3 .	.34	3.03	1.81	0.36	3.03	4
4	755276	AA498349	0.11	0.30	2.74	3.5/	4.02	*	14.34	134.4	1.13	ဂ
8	270917	N32514	0.01	0.23	25.82	¥.	5.1.	2 ;	0.61	3.63	22.02	က
15		AA431721	0.20	3.10	15.42	64.92	77.	0 0	71.36	3.78	2.68	က
152	320485	W16659	90.0	0.88	13.89	1.39	3.75	š !	78.47	44 85	2.05	ო
5	293579	N94111	96.0	12.90	13.42	0.73	1.52	77.4	14:07	030	1.22	ю
2	281039	N47717	0.78	8.8	12.85	43.16	25.22	F0.9	77.7	48.34	0.82	က
. A		AA116781	20.0	0.48	12.37	48.34	8.67	1.53	77.0		3.44	က
3 2		7316744	0.08	0.92	11.12	0.12	0.12	2.23	99.00		1.67	n
3 5		UKO1KB	190	5,43	10.69	20.89	31.93	7.51	1.20	9 7		· e
2 5		NA 8250	180	18.34	10.17	21.89	29.40	4.51	<b>4</b> .88	0.0	2 7	) e7
3 5		44448877	807	33.97	8.32	14.84	26.99	1.81	1.28	0 6	9 6	
2		100000	90.	23.62	8.02	1.97	4.75	1.05	5.38	32.20	700	•
3	470610	A A C 20 B K 4	20.0	0.57	7.75	3.54	3.42	1.63	1.38	34.08	7 7 7	n e
9		20000	;	ğ	7.33	0.87	3.83	1.16	32.30	5.68	7 (	, (
<u> </u>		1.1CB+N	2 5	, ,	7 13	4.67	30.78	1.10	3.88	66.0	1.36	<b>n</b> (
豆		AA43/241	0.4	5.5	9	2.41	4.29	2.60	21.59	8.58	1.59	n
Ž	•	AA040861	9.20	80.00	3 6	13 15	16.83	1.72	1.58	1.47	3.24	n .
इ		N74131	0.0		200	18	2.98	4.25	20.28	2.26	0.61	r)
8		AA233809	50.0	57.0	47.9	1 70	4.23	0.65	4.75	0.56	22.68	ო
167	757389	AA437142	0.02	<u>-</u>		• !	ų					
							c					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

		Ana Hank			Fold(MPM123/A Fold(MPM124/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A # of Tumor	A)0019761877	End (MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A F	old(MPM126/A	# of Tumor
	MAGE Gode D	Accession	AVEBPH	AveTumor	Fold(Ave I umoriav •BPH)	veBPH)	vaBPH)	veBPH)	veBPH)	veBPH)	veBPH)	Samples
		Number	1	3	9	171	1.65	2.18	16.65	4.09	7.3	- e
ş	196168	R92362	0.59	3.32	00.6	- 00	18.59	4.47	1.32	1.05	2.26	n
169	687867	AA235370	0.81	<b>3</b>	BC 16		14.24	8.62	0.72	0.88	2.12	က
5	712950	AA282273	5.93	32.98	9:30	. e	0.51	7.72	0.13	17.97	0.10	ю
Ē	504761	AA148735	0.08		o	9.50	₹08	0.73	20.36	0.89	2.62	m
Ē	32243	W16424	0.23	<u> </u>	9.30	. i.	4.65	2.60	1.65	11.48	99.9	60
E	35010	R45118	0.66	53	6.03	4.14 2.63 2.63	2.75	808	66.0	7.48	8.28	က
17.	762537	AA418484	1.72	9.60	5.01	70.7	23.43	2 17	08.0	1.13	5.84	n
175	277506	N57002	1.48	7.7	4.97	99.7	1 00	: Z	12.13	2.89	3.52	က
176	773208	AA425650	9.71	47.00	4.85	1.82	08.7	13 33	96	7.22	4.39	6
Ħ	364865	AA035745	0.44	2.09	4.78	1.52	1.24	1 64	4.85	4	0.89	69
178	247587	N58136	0.01	0.06	¥.	1.96	14.28	<u>.</u>	3 8	1.68	2.65	n
179	34888	R19878	0.14	0.67	4.62	5.69	9.05	4	2 .	4.55	0.13	6
8	72385	T51639	0.08	0.36	4.59	0.67	18.34	4.47	5.5	2.45	3.58	n
181	773204	AA426892	12.74	58.08	4.56	1.47	1.49	5.37	13.00	0 F. T	7.68	. 67
182	277327	N57483	0.02	0.09	4.58	1.67	6.21	2.56	8.18	6. 6	9 6	
3	770088	AA430545	7.38	33.38	4.52	7.28	8.45	8.19	<b>X</b>	0.26	00.7	, «
3 3	51820	HIDEES	0.03	0.12	4.40	1.90	8.33	7.29	6.63	2.43	0.30	, ,
<u> </u>	206347	S S S S S S S S S S S S S S S S S S S	0.70	3.10	7.	8.62	98.8	4.53	<b>79</b> .0	0.94	1.83	n (
3 5		745020		1.67	4.40	4.09	12.69	<b>8</b> .8	0.91	1.09	1.27	י ניי
8 9	7,1087	W02420	? .		. ₹	0.98	7.98	1.80	5.23	1.24	<b>8</b> .83	ro -
18/	1009/7	851004		100	7.33	3.30	1.67	0.98	9.74	9.64	9.67	က
<b>8</b>	22000/	AA128407	70.0	5 6	4.33	10.35	0.18	0.28	2.13	8.96	<del>1.1</del>	n
<b>2</b>	97560	<b>74545</b> /0	8 8	97.0		0.46	1.86	0.68	3.72	15.21	3.60	n
<u>\$</u>	202058	8	60.0	7 :		7.30	9.78	3.67	2.18	1.01	1.74	<b>6</b>
<u>=</u>	285997	N73563	0.42	1.61	07:4	4 5 6	2.40	5.57	1.28	1.92	8.78	9
5	205828	H57585	LC:0	61.7	07.4	90:-	68.2	0.68	0.90	0.68	9.45	က
5	67397	149325	10.0 10.0	8 5	5.10	3.45	0.81	2.68	7.58	1.37	8.82	က
<u>\$</u>	811028	AA465373	8 3	20.42	4 6	2 07	1.98	7.47	8.01	2.44	1.39	n
<b>5</b>	489637	AA101878	7.80	79.	9. 6	500	61.6	6.46	2.24	5.69	1.52	က
<b>8</b>	284624	NS4790	 	9.50	<b>5</b> 0°+	9	03.6	4.10	3.36	1.15	208	<b>6</b>
187	641507	AA487385	<del>.</del>	87.7	70.4	0.38	7.83	0.28	9.35	0.79	5.34	6
108	882407	AA676604	<b>5</b>	41.0	\$ (	6.50	90.0	1	2.63	4.98	<b>4</b> .00	n
<u>\$</u>	280155	N#7012	8	0.25	3.82	2.32	77.	A 78	260	1.68	2.22	က
<b>50</b>	1466788	AA883735	0.46	1.70	3.80	2.83	r 4.0	2	88	1.93	11.30	6
201	784589	<b>AA44</b> 3300	0.03	0.10	3.89	3.05	66.1	t	2			
							9					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	Mumber	AVEBPH	AveTumor	eBPH) veBPH) veBPH) veBPH) veBPH) veBPH) veBPH)	veBPH)	veBPH)	veBPH)	veBPH)	veBPH)	veBPH)	Samples
278877	AA202583	0.01	0.05	3.80	3.91	7.78	1.48	8.70	₽2.0	9.74	က
858174	AA630620	0.03	0.11	3.69	6.41	6.18	98.0	0.35	1.33	8.10	n
782575	4447522	0.38	1.39	3.86	8.55	7.01	2.57	0.65	0.88	3.47	က
428166	AA001870	2.28	8.77	3.85	1.91	3.10	15.	10.12	1.60	1.78	က
588828	AA130857	1.17	4.40	3.83	2.76	2.15	2.16	8.17	4:00	3.75	က
730838	A 4417012	0.02	0.00	3.61	7.50	0.49	0.40	7.80	2.28	<b>4</b> .30	60
837053	44468478	7.85	29.78	3.79	3.76	5.83	1.10	2.48	6.86	2.82	ო
K07824	A & 4.3.3050	22	1	3.78	1,62	60.0	1.78	6.89	6.74	6.57	က
208210	HASSA	0.89	335	3.76	5.28	5.92	7.33	1.69	0.78	1.61	က
427996	AA001841	110	0.43	3.75	3.08	1.05	3.00	3.88	0.51	11.00	6
123755	P01304	28	989	3.75	0.50	0.29	0.34	4.24	12.61	4.50	က
436153	AA703249	14.14	52.21	3.69	1.02	1.89	3.14	8.23	5.06	2.80	ო
143062	R71335	770	1.61	3.67	2.83	9.9	2.57	2.67	3.32	4.58	က
131599	707FCA	0.16	09.0	3.65	5.05	2.39	2.70	1.07	4.57	8.14	<b>6</b>
240151	H82706	2.33	8.40	3.64	2.83	9.0	1.92	4.91	6.33	5.22	es
452118	AA707219	4.0	1.61	3.63	2.41	7.49	3.46	2.29	3.62	2.52	<b>~</b>
201125	R98472	1.05	3.81	3.61	3.83	4.00	2.14	6.14	2.91	2.55	က
126713	R07115	0.27	0.98	3.55	6.64	4.18	2.20	4.29	1.20	2.78	က
768179	AA424849	2.56	90.6	3.54	3.04	3.99	1.69	2.78	7.17	2.81	က
250678	H95976	60.4	14.42	3.52	2.85	6.36	3.23	3.77	2.33	2.51	က
273499	N33263	80	0.32	3.52	3.88	2.67	2.75	6.21	4.11	1.47	n
765765	AA486455	1.95	48.9	3.51	2.12	3.80	3.48	7.92	<u>1</u> .92	1.82	eo
815047	AA485168	2.68	9.3 X	3.48	2.62	1.60	1.82	4.71	3.01	7.18	<b>6</b>
384670	AA709333	4.17	14.53	3.48	3.11	2.41	3.77	7.25	2.34	2.03	က
782200	AA431964	1.53	5.30	3.47	2.48	6.32	2.54	3.94	2.26	3.20	ы
52730	HZ8500	<b>8</b>	22.94	3.45	2.33	2.11	4.07	6.25	3.48	2.49	ო
125311	R05810	1.82	6.62	3.6	1.07	2.84	3.69	6.68	4.33	2.05	က
770880	AA434404	8.05	27.88	3.44	6.26	3.15	7.87	1.10	0.49	1.77	ဂ
796268	AA480827	0.55	1.89	3.43	1.43	3.68	2.58	5.27	5.89	1.7	ы
204148	H55920	2.07	7.07	3.42	2.91	3.75	7.21	4.16	1.12	1.39	က
811373	AA178833	0.01	9.0	3.41	9.82	3.38	1.53	3.86	0.80	0.90	က
325380	AA284285	0.04	0.03	3.40	1.86	1.13	5.78	5.41	1.13	5.09	<b>6</b>
434822	AA703114	0.80	2.72	3.38	5.38	1.97	4.32	1.12	1.58	6.01	က
1040033											

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

		Gen Bank			ERIMANATIONAL FOILIMPM201A FOICIMPM121/A FOICIMPM124/A FOICIMPM122/A FOICIMPM123/A FOICIMPM126/A \$ Of Tumor	Fold/MPM120/A F	old(MPM121/A F	old(MPM124/A F	old (MPM122/A	Fold[MPM123/A F	old(MPM12E/A	# of Tumor
	Clone ID	Accession	AveBPH	AveTumor	(BPH)	veBPH)	veBPH)	veBPH)	veBPH)	veBPH)	veBPH)	end Has
;			,	ě	o c	75	4 86	2.99	5.61	3.78	1.72	n
238	882483	AA676698	2.75	9	3.30	<u> </u>	72.0	1.14	3.10	10.17	4.12	ო
237	767130	AA424535	0.40	1.67	3.38	3. 6	7 7	1	7.41	0.38	2.21	က
នី	126321	R06479	8	0.10	3.38	71.7		. 77.	2.89	8.21	0.15	ဗ
239	230440	H75328	0.07	0.22	3.37	1.97	3.37	, c	203	2.93	1.89	က
240	624271	AA181179	8.78	22.81	3.36	3.53	3.83	70.7	4 6	95	0.58	6
241	123265	R00284	2.30	7.72	3.35	4.22	9.13	4.42	0.68	28.5	<b>4</b>	
242	357238	W93838	0.72	2.40	3.35	5.78	5.69	1.43	1.48	3.53	<u> </u>	,
243	122331	B00479	8	3.43	3.33	2.37	4.72	99'9	98.0	3.48	7.50	, c
77.	240404	728437	8	6.17	3.33	2.70	3.07	1.59	57.7	3.58	1.21	
	347482	******	3 2	6	3.32	0.45	5.03	5.53	4.94	2.81	1.19	r <b>9</b>
	2000	110001	3 8	; <del>;</del>	334	3	4.36	6.28	4.13	2.87	0.31	n
240	28020	HZ3Z3Z	3 ;	- 8		 200	53	3.17	4.45	2.72	2.93	m
247	160628	H25019	<b>1</b>	5 6	3.30	70.7	2.43	7.21	1.55	3.28	3.34	n
248	1505294	AA905838	1.1	3.68	3.30	O8: - 6	2.43	3.49	6.75	1.98	1.68	က
249	262053	H98812	7.69	25.28	3.28	8.7	2+;c	3.14	25.	2.25	5.58	က
250	741067	AA478436	17.58	57.76	3.28	2.50	9.59	2 6	4.85	1.83	3.72	m
251	490178	AA121266	3.50	11.46	3.28	2.64	¥.	077	0. 6	2 83	127	m
252	208741	H63175	0.09	0.30	3.26	2:00	4.61	3.24	40.7	70.7	20.6	•
253	416811	W86466	3.63	11.85	3.26	2.01	2.14	2.13	5.38	4.80	3.0	, ,
25.	418082	W85881	05.0	1.62	3.26	2:00	2.31	1.93	3.05	3.55	6.62	, c
255	221581	H92525	0.55	2.	3.26	1.18	1.35	3.60	1.43	<b>9</b> .	5.05	י כי
3 5	22.274	LABAKIA	50	0.17	3.25	3.24	0.10	0.19	2.95	5.14	7.79	n .
3 5	46287	2000		* 2	3.24	8.67	5.69	4.15	1.74	3.08	1.09	n
ē i	1000	20000	. v	9 6	3.23	2.48	3.52	2.19	3.80	2.05	6.34	ED.
8 5	70/011	ACCO.	6.4	3 5	3.23	1.59	4.12	2.42	5.41	3.80	2.03	•
RC7	044171	70//81	- 6	2. C	322	4.87	5.06	1.22	2.93	3.17	2.10	e
3 3	710701	1678	3 6		3.22	08.	<b>+</b> .01	2.67	3.29	2.89	4.45	n
		91///	5 5	2 01	32.	2.30	5.27	1.85	3.83	2.74	3.25	<b>~</b>
70 6	2000	2000	; ;	8	3.21	1.59	2.24	3.47	5.80	5.20	0.85	n
3	0.000 A	/8001	, a		9 t 6	88.	1.58	3.78	3.20	2.11	6.82	6
\$	8	10406	9 9	2 5		08.7	29	2.43	2.00	3.28	2.91	n
582	4/02/4	WBGZUZ	2 G	2.5	, c	3.60	3.81	2.60	5.88	1.06	<b>3</b> .08	c
90 1	1422/1	AA62/405	7 6	? 6		27.	4 27	1.59	8.08	3.23	0.45	c
/97	13/01/	6000	3 6	8 8		66.6	3.85	2.53	3.68	3.77	2.82	n
897	459/97 4500	711800	7.0	8 5	5. c	1.18	4.57	2.43	1.60	1.72	₹: •	e
802	437537	/ZMRRGWV	0.0	<b>8</b>	<u>.</u>			i				
						_	89					

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Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	IMAGE	Gen Bank Accession	AveBPH	AveTumor	Fold(AveTumoríAv Fold(MPM120/A Fold(MPM121/A Fold(MPM122/A Fold(MPM122/A Fold(MPM126/A \$ of Tumor ARPH) veBPH) veBPH) VeBPH) veBPH) VeBPH) VeBPH) VeBPH) VeBPH)	Fold(MPM120/A	Fold(MPM121/A veBPH)	Fold(MPM124/A vebph)	Fold(MPM122/A veBPH)	Fold(MPM123/A veBPH)	Fold(MPM125// veBPH)	Samples
		Number						76.0	. 33	2.73	4.79	က
270	1475734	AA872692	7.38	23.26	9.15 5	3.45	3.20	<b>1</b> 0.7	40.4	) o		٣
27.1	268876	N26026	0.61	1.59	3.14	1.46	6.62	3.35	1.38	6.00	6.16	,
272	79032	T61899	16.17	47.49	3.13	3.66	4.92	3.98	1.33	2.11	7.80	, (
773	771023	AA427978	1.12	3.51	3.13	2.45	3.57	2.85	2.35	3.41	4.13	m
2 2		A 4 4 8 5 8 7 8	2.17	27.5	3.09	9.1	2.45	1.78	5.54	4.36	3.37	m
7	24044		: g	25	60 6	76	3.31	2.15	127	5.53	2.34	es
67.5	1710/7	A A 702/80	9 6	, «	80.6	3.50	2.77	3.84	5.73	1.38	127	က
9 1	71.004	AA 100557	3 6	2 47	3 07	2.06	2.12	3.31	5.09	3.43	2.39	က
7 6	10001	A & 80 28 24	5 C	2 2	3.05	166	4.32	2.32	3.80	1.77	4.45	es
9 6	768746	134070	6 6	5.87	305	2.21	3.82	1.69	1.33	5.44	3.82	60
	170303	7471044	9 0	55	100	<b>4</b> .39	4.19	1.6	1.95	2.28	3.78	က
	80050	74224F	12.72	38.68	<b>1</b> 0	10.1	1.73	3.23	4.87	5.48	1.91	60
	4500182	A A 998720	8 1	5.73	70 6	2.12	4.97	127	4.24	1.94	3.69	က
787	288724	N20007	2	5.20	3.03	2.45	3.22	6.41	2.32	1.71	3.08	က
3 2	100623	DORFRE	0.51	35	3.03	2.89	4.10	3.93	3.02	2.70	<del>1.</del>	60
, K	41205	PARR77	19.83	80.07	3.03	2.41	3.80	2.88	4.40	3.55	1.08	•
286	795555	AA459879	8	ā	3.03	3.06	5.11	2.63	4.91	1.14	132	<b>6</b>
287	416808	W/86779	0.57	1.7	3.02	1.77	4.00	2.37	4.02	3.51	2.42	က
288	242698	H94282	1.16	3.50	3.00	1.70	2.11	3.21	4.15	4.88	1.97	က
289	364352	AA022498	0.03	0.08	3.00	2.52	0.38	4.22	3.89	4.83	2.18	ന
88	244050	N34042	90.0	0.17	3.00	1.60	3.79	1.68	2.22	3.29	5.43	က
291	544639	AA074677	1.05	3.15	2.99	1.72	3.77	3.39	3.80	2.68	2.58	m
202	862520	AA663058	17.95	53.54	2.98	3.39	3.07	2.81	4.89	2.83	0.82	m
283	797000	AA463500	0.11	0.33	2.98	<b>1</b> 5.1	3.21	1.65	5.62	1.07	4.77	(m)
787	1483252	AA886870	18.18	54.08	2.98	4.82	3.25	2.81	3.28	1.52	2.20	က
295	447416	AA702335	1.15	3.41	2.96	5.58	4.11	2.12	40.1	3.10	<del>2</del> .	e
296	1475308	AA877997	2	4.56	2.86	2.82	3.27	4.53	1.90	1.03	3.19	ო
297	1212	H51039	0.22	0.85	2.95	1.35	3.39	3.88	2.74	3.89	2.47	<u>ب</u>
298	1456937	AA863449	1.15	3.38	2.84	5.57	3.38	1.71	2.34	0.80	3.82	က
288	201039	H48278	0.83	2.42	2.93	2.76	3.98	2.31	3.08	2.04	3.43	က
300	263769	H99894	0.75	2.19	2.83	3.82	7.88	3.41	1.10	1.06	0.29	က
301	427830	AA001976	0.05	41.0	2.91	0.99	3.14	6.49	0.21	7.7	2.22	က
302	307244	N95187	0.35	1.01	2.91	2.20	4.92	3.17	2.07	3.85	1.25	က
8	810843	AA458959	90.0	0.17	2.90	134	3.15	1.41	5.84	177	<del>1</del> .	က
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Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	MAGE	Gen Bank			Foldiave Tumoriay Foldinphitzia Foldimphitzia Foldimphitzia Foldinphitzia Foldinphitzia Foldinphitzia # of Tumor	Fold(MPM120/A	Fold(MPM121/A	Fold (MPN124/A	FOICH MPM122/A	Fold(MPM123/A	Fold(MPM125/)	a of Tumor
	Clone ID	Accession	AveBPH	AveTumor	•BPH)	VeBPH)	veBPH)	veBPH)	VeBPH)	VeBPH)	(W-88-W)	e adupa
Š	000001	100000 V	183	10.40	2.80	2.35	2.11	3.29	1.57	3.43	4.63	n
5 8	1400020	AA888/83	20.0	B	08.6	3.08	3.41	3.69	2.10	2.31	2.77	က
3	AC707	145024	ò 6	8 8	9 6	3.87	3.77	1.00	1.23	1.00	6.23	ი
3	X	AA062813	5.0	3 6	6.07		4.75	3.79	1.52	0.39	3.84	က
304	262104	N51498	5 6 6	6.00	2.0.4 18.6	3.18	3.47	5.50	1.78	1.21	1.74	က
	344308	W70147	9 6	<u> </u>	2.01	2.0	3.24	8. E.	3.52	2.33	2.40	က
8	6266	H06725	2.3	<u>.</u>	2.6 Ce.	2 7.8	4.53	306	2.48	3.30	0.85	၈
310	196303	R92435	<b>3</b> 5	<u> </u>	7.50	99.5	3.24	2.59	3.41	1.52	2.19	ო
E :	207483	N25262	<b>8</b> 0.	<b>?</b> ;	11.7	2 6	187	3.45	1.24	1.90	1.53	က
342	757198	AA496133	0.65	87.1	2.73	5. c.	4.23	3.40	3.07	2.93	0.52	n
343	204122	H81886	F	n i	6.73	6.30	60 0	38	1.18	3.64	3.82	n
314	839545	A489791	0.42	<b>7</b>	2.73	7	3 9	0.6	2.89	4.45	4.47	ь
315	809694	<b>22.7</b> 22	0.02	600	2.70	<b>,</b> 5	0.50	231	1.13	3.56	3.32	6
316	126234	R06362	0.30	0.61	A0:7	2		10.	3.28	3.22	2.72	က
317	280407	N51583	0.87	2.62	2.69		0 F C		3.48	3.11	4.17	en
318	758284	AA404231	0.78	2.08	79.7	707	00.6	3.23	204	1.07	2.08	n
310	150314	H00817	2.18	5.61	7.67	9 0° 7	3.47	3.00	88.0	2.99	3.87	e
ន្ត	288936	N62698	3 :	0.11	7.07		,	1 79	55	0.21	5.08	e.
33	767440	AA437228	0.03	0.13	2.67	0.27	. t	 	2.5	80	1.30	r
g	178554	H45289	0.87	2.28	7.07	1.08	2.02	1 2	+ BK	3.26	3.01	ო
323	288057	N69989	0.68	1.79	2.62	2.03	3.01	¥ .	69:	0.70	386	, m
88	755299	AA496359	1.60	4.39	2.80	4.47	0.52			162	57	e
32	725321	AA291749	0.87	2.27	2.59	3.21	3.07	3.01	3 5	3.58	4.52	. 63
g	1467831	AA883431	1.17	3.02	2.59	1.16	77.1	7.35	4 . C	3 04	334	· 67
321	120097	T95151	0.60	<b>3</b> 5	2.58	2.09	2.07	<u>•</u> ;	9.0		3	
328	139455	H15746	0.02	0. \$	2.58	0.62	3.70	231	. c.	3.72	23	· •
8	1460698	AAB&8802	6.57	16.82	2.58	1.39	3.08	<u>\$</u>	7		1 8	• •
93	757368	AA437128	0.02	90:0	2.55	3.92	2.20	3.37	0.77	0.24	3	n (
331	450612	AA682452	0.74	1.89	2.65	1.17	96.0	0.92	3.00	4.79	1	m (
332	342543	W88141	0.05	9.0	2.53	4.17	1.	0.58	1.17	4.50	0.30 0.30	m
333	503741	AA131486	1.05	2.05	2.53	3.31	3.85	3.58	1.37	1.10	<del>1</del> 88	ю
ğ	784130	AA432075	57.71	44.77	2.52	3.17	4.60	0.78	1.58	3.28	1.79	က
33	325370	W52208	0.03	0.08	2.51	3.68	3.87	2.82	1.13	0.31		ဗ
82	288663	N62394	0.76	1.95	2.50	3.07	3.86	0.85	<b>7</b> .00	1.37	3.87	n
B	344359	W73377	0.07	0.18	2.46	0.90	3.52	0.14	3.08	7.02	9.08	က
							10					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

		Gen Bank		•		Enid(MPM120/A	Fold/MPM121/A	Fold(MPM124/A	Fold(MPM122/A	FELLINGTON ENGINE FOLDING POLICIES FOR PRINCE FOR FOLDING FOLDING FOLDING FOLDING FOLDING STEELS FOLDING	Fold(MrM125/A	A # of Tumor
	IMAGE Clone ID	Accession	AveBPH	AveTumor	Fold(AVE) GIRDINA •BPH)	VeBPH)	veBPH)	veBPH)	veBPH)	veBPH)	VOBPH)	e and iller
		Number			. :	ć	7.9 5.	76.6	1.40	1.23	1.35	n
338	63381	R16241	1.15	2.79	2.41	3.60	6.0	5 5	1 85	1.08	3.55	က
339	429823	AA009803	0.4	0.35	2.40	3.16	3.19	70.	9	40.0	232	ю
35	258242	N20855	0.62	1.49	2.40	3.61	3.23	0.28	4.12	5.0		
3		90000	000	0.00	2.31	0.60	4.21	<b>1</b> .00	1.52	3.40	• ·	•
3	/89CCZ	NZ/308	9 6 0	2 6	5	128	3.24	1.70	3.10	3.01	1.39	<b>.</b>
342	452068	A707125	2 2 2 3	3 3	7.30 30.00	64.6	3.80	0.16	3.57	0.38	3.18	n
35	565778	AA135870	20.69	48.84 48.64	27.7	2 G	2 6	0.24	3.30	0.24	<del>4</del> .05	က
317	128508	R10154	<b>5</b> 0.0	0.08	2.15	07.1	5 6		20	0.88	0.82	က
345	839048	AA487505	0.36	97.0	2.12	4.12	6.45 6.45	7 6	 	4.85	73.63	8
348	280763	N50583	0.03	0.50	14.38	0.58	2.77	2,38	97.	13.48	32.88	2
772	FDOR 3	AA158182	0.01	0.18	13.25	0.74	0.74	<del>-</del>	<b>4</b> :	04:34 0:30		
Š 3	200	2000	6	0.22	11.30	1.40	0.82	3.37	1.36	2.48	58.40	4 (
8	5	AA620472	70.0	4 ;	8 9	1.48	2.83	2.36	49.00	1.07	331	8
3	377610	AA056833	0.13	7. T	10:02	0 F	97.0	2.38	2.67	34.78	5.82	8
320	50787	H16833	<b>0</b> 0	0.50	2.80	90.1	2 8	750	0.56	1.1	2,44	8
351	137984	R63085	0.29	2.25	7.79	8.43	33.82	5		27 14	1.71	84
352	858585	AA669222	0.12	0.8	7.77	1,46	0.70	9.1.	٠ ٢	7	80.0	•
24.2	358000	W872#3	0.08	0.49	7.64	1.97	2.74	4.55	0.24	39.70	8 6	4 C
3 3		5077844	5	0.75	7.47	33.12	2.43	1.18	3.78	1.47	<b>5</b> .7	7 (
Š	430192	AAUTUSZB	2 (	9 0	100	7.32	29.35	1.05	0.49	0.48	2.55	7
355	151477	H02837	B !	0.10	0.0	46.33	19.31	2.50	1.6	0.68	1.40	7
358	81331	T60111	0.47	3.18	ğ.	5.55		130	1.63	1.01	90.38	7
357	52026	H22832	0.25	1.66	6.69	13.67	99.77	, c	2 2	11.38	5.36	8
88	859627	AA686405	0.08	0.53	6.43	1.07	0.12	0.12	20:0	9 67	18 47	0
250	4200RB	A A DOS 198	0.22	1.38	6.27	0.88	0.41	0.38	<u>د</u>	04.0		
3 3		104033	8	0.32	6.48	6.98	2.35	1.09	0.55	1.30	20.60	•
3		25.55	, ,	17 OB	979	2.04	2.77	1.7	2.77	11.30	12.03	<b>.</b> 4
Ş	00540/	AV430142	3 6	23.4		1 16	22.31	1.49	3.68	0.91	.58 .58	7
88	788226	AA424900	8 8	5.0		4 05	23.41	98.0	1,45	0.13	0.89	8
8	303138	N90808	70.0	1/7	- 6	27.0	151	78.0	20.25	2.06	5.30	8
Ř	757165	AA443850	0.0	D	3. C	;	7.7	2 52	1.47	0.61	2.40	7
365	565235	AA136125	18.25	89.08	88.4	00.7	2 .	18.	0.22	2.88	18.58	7
8	247103	N53858	0.05	0.22	4.82	9.79	<u> </u>		6	72.71	2.86	8
387	234378	N28268	7.87	14.16	4.81	1.46	2.58	4.82	70.7	<u> </u>	3 6	
38.	758405	A4482119	9	0,21	4.73	0.22	1.45	2.24	3.15	2.40	9.6	۷ (
8 8	47755	H10081	0.10	0.48	4.65	14.30	0.10	0.55	0.22	9.85	2.80	<b>79</b>
3 8	2002	000	076	15.28	4.38	5.72	2.44	11.68	129	2.30	23 28	7
	37.000	080//100	7 7	1 05	4.33	6.41	14.50	1.23	1.41	1.47	<b>0</b> .83	7
5	7/004/	AA020213	?	2	•							
							=					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

		Gen Benk				A A TOCARDONAL STATE OF THE STA	Pold/MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A F	old(MPM126/A	# of Tumor
	Clore 1D	Accession	AveBPH	AveTumor	FOIG(AVE) UIROUAV •BPH)	VeBPH)	veBPH)	veBPH)	VeBPH)	veBPH)	veBPH)	Samples
		Number 1	:	į	Š		7.49	1.07	1.48	90:0	19.71	7
22	29782	T63861	0.1 <b>8</b>	0.71	4.32	- !		900	1 70	0.83	4.17	7
S S	420424	AA007887	0.03	0.14	4.20	2.13	00.01	2 10		\$	0.11	2
374	265855	N22836	0.0	0.39	4.17	16.98	1.24	1.97		3 -	1.42	۰
175	1808315	AA001108	0.72	3.01	4.18	2.17	2.93	3.20	13.79	Į.	7	• (
3 6		2 4 9 5 6 4 6 0	, Y	0.00	411	11.10	10.86	0.50	0.52	0.21	<del>S</del>	7
5	0108001	90(009A)	3 .	3 6	7 00	1.22	10.53	5.68	2.67	1.88	<b>2</b>	8
3/	CSS323	4	2 6	4 5	9 6	2.68	8.38	0.83	0.57	0.82	10.67	7
378	284831	N71861	0.02	) (	<b>a</b> 6		0.36	5.13	2.	0.36	15.08	7
379	1031203	AA809982	0.03	C	0.°C	71.7	0.0	2.18	4.17	0.30	16.49	8
98 98	758360	AA404280	0.03	0.13	3.05	0.30	000	2 G	2.97	1.40	10.58	7
ž	178569	H49053	0.05	0.20	3.85	3.10	9 1	9	1.67	2.16	12.53	7
382	164477	R54855	1.01	3.95	3.92	2.32	c/:l	2 6	5 8	£	95	~
88	1048998	AA778636	0.14	0.57	3.91	10.73	5.96	2.86	77 7	S	8 5	. ~
385	129589	R16541	4.0	1.58	3.79	2.77	13.43	1.08	0.83	0 ;	B 1	4 6
7 8	10KRA7R	A A R 20027	91.0	0.72	3.75	3.32	2.92	2.58	272	1.71	17.8	<b>,</b>
3 8	484800	44682274	10.27	38.51	3.75	0.81	2.58	2.98	8.45	6.06	<u>2</u> .	74
8 8	8010	*/7700V			3.74	1.87	2.97	2.88	2.21	5.65	6.87	7
3	78006	070/14	2 6	2 5	7.6	2.76	5.27	1.60	11.37	0.71	0.71	7
9	078706	AAUTOBOU	<u>}</u>	3 3	- 6	0.66	8.21	25.0	<u>z</u> .	0.20	10.81	7
8	122178	198615	20.0	C 6	<b>9</b> 0.0	0.00 4.50 4.50 4.50 4.50 4.50 4.50 4.50	12.29	1.20	28.0	0.62	0.85	74
8	246808	N53214	20.5	8.7	20.0		8 7 A	2.83	2.98	5.39	1.99	7
8	194888	H40350	0.61	2.24	3.68	5. 5. 5. 5.	9 8	2.18	0.92	0.72	1.33	~
302	155072	R71383	0.30	1.07	3.62	00.7	0.3	2 6		7.54	3.33	7
383	190325	H29897	0.70	2.51	3.59	2.78	2.7	10.7	7 6 6	4 62	6.75	7
ğ	502438	AA134862	1.10	<b>1</b> .0	3.58	1.28	2.12	2.17	70.7	*O'.	) e	
385	283689	N69675	0.0	0.03	3.57	6.95	1.08	1.08	8.	10.16	0.T	•
8	1486752	AA912071	0.70	2.48	3.57	1.94	3.07	2.18	10.40	1.10	2.70	7
į	840438	44053220	0 03	0.10	3.52	6.17	0.37	7.88	1.98	1.85	2.87	7
	7804.5	4487020	2.86	2	3.45	1.42	1.18	2.71	7.18	5.88	2.39	7
		Nanna.	2	2	3.45	3.79	2.25	7.49	2,62	2.54	1.99	. 7
3 (	0/4170	W32303	3 ?	20.4	7.	2.88	4.89	2.73	1.12	7.17	1.77	~
3	000/07	1402332	<u> </u>		97	2.43	3,66	2.76	7.36	2.52	1.53	7
5	256300	2000	2 6	7.7.	00.0	9 9	27.6	3.67	2.20	2.53	6.91	~
ğ i	1048040	AA000043	78.0	3 5	30	2 6	6.28	181	6.92	1.89	1.88	8
\$	289025	N69826	0.00	797	cc.e	7.7			2 13	1.50	1.49	7
\$	291708	N73480	0.53	1.7	3.33	4:14	1.81	9.8	 	1.62	4.45	
405	1031568	AA609304	0.08	0.25	3.30	6.38	2.97	1.77	19:7	70.1	ř	•
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Table 2-1;T3NO vs. BPH; Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

		7000		'		Alocation	Ecid(MPM124/A	Fold/MPM124/A	Fold[MPM122/A		old (MPM126/A	# of Tumor
	Close ID	Accession	AveBPH	AveTumor	old(Ave lumona) eBPH)	veBPH)	veBPH)	VeBPH)	veBPH)	veBPH)	veBPH)	Samples
		Number				1	308	2.77	2.09	2.39	1.81	7
8	383676	AA679116	0.13	0.41	3.28	66.7	5 5	2.24	4.55	5.37	2.84	7
407	310483	W31074	6.22	20.37	3.28	2.57	2.10 4.10	2 97	2.78	0.15	0.89	8
408	84713	174257	0.0	0.21	3.27	6.85	0 0 0 1	1 2	12.99	* <del>*</del>	0.35	8
409	219955	H81817	0.18	0.61	3.24	0.75	0.51	9.10	12.6	4.35	6.93	2
410	295741	W02483	0.35	1.13	3.23	20,	2.14	8 8	. a	1.58	5.09	7
<b>1</b>	796848	AA460115	9.12	28.44	3,23	1.38	0.93	A0.	4 E	1.48	10.68	2
412	44253	H05818	90.0	0.21	3.23	<u>.</u>	1.21	0.62	9 6	3.85	2.88	7
413	139051	R62828	0. 4	1.43	3.23	2.91	2.95	4.57	80.2 0 50	00.6	1.53	2
414	454543	AA677023	0.83	2.67	3.22	0.89	1.39	2.00	10.50	20.0	225	2
***	40316	H16368	0.22	0.71	3.21	69:0	9.1	<u>4</u> .	9:50	7.12	983	۰ ،
2 4	348412	W74482	1.22	3.89	3.17	2.96	9.86	0.85	1.50	3.24	700	٠,
} ;	283838	H00788	1	3.52	3.17	2.49	4.47	2.58	8.53	1.43	SG- 0	٠,
;	200000	0000000		2.2R	6. 2.	7.40	1.58	1.99	4.43	2.52	0.00	7 (
5	28/0/2	200000000000000000000000000000000000000	# F	9 6	3.13	0.71	1.64	1.65	7.85	<b>₹</b> .	2.01	NI (
418	71987	C/97CN	9 4	9 7	3 13	5.71	7.10	2.75	1.53	0.65	<del>.</del> .08	2
450	1084274	A1003028	÷ ;	, u		2.46	2.18	2.50	3.58	6.73	1.25	~
421	784212	AA446864	[R: 0	G .	; ;	96:0	3.88	1.59	2.55	0.17	9.50	~
<b>4</b> 22	41214	K26898	0.0°	- 6	- 6	7.63	1.12	4.50	2.52	0.37	2.48	7
423	361688	W96197	0.03	8 5 6	2 6	5 5	107	3.65	8.67	2.39	2.00	7
454	148013	R82299	7.60	73.51	3.08	. 27.6	700	2.83	2.19	4.07	4,03	~
425	392388	AA707935	0.52	1.58	3.09	2.30	6.0	98	3.28	0.17	0.99	7
450	347615	W81526	0.07	0.20	3.08	12.89	0.30	55.	1.66	4.08	8.48	7
427	1030779	AA609004	0.52	1.61	3.08	1.28	o e	84	133	5.86	1.60	7
428	384555	AA022601	<b>Z</b> .	5.67	3.07	5.3b	6.13	3 6	5.17	1.89	1.28	2
429	743688	AA829324	1.27	3.80	3.07	2.60	17.7	07:0	. u	9	0.24	8
430	141726	R69584	0.0	0.13	3.07	8.46	2.87	3.48	2 .	F 7	4.82	8
5	796147	AA460988	2.96	90.6	3.07	2.49	2.65	2.38	<u> </u>	5 6	5.35	2
432	813501	AA456078	0.60	1.85	3.07	1.76	2.60	4.83	7.72	5 5	88 7	۰ ،
13	138047	853757	0.35	1.07	3.05	2.90	1.81	2.18	2.33	4.18	8 :	4 (
3	22800	TA7236	0.12	0.37	300	3.86	5.88	1.88	2.82	2.48	C :	<b>7</b>
} ;	20484		9	0.48	70.6	2.44	5.47	4.57	1.88	2.74	1.14	7
6	248784	1472074	200	2.80	70 6	2.94	5.24	1.48	2.88	3.18	2.50	64
2 5	10/05	170774	,	78.0	3.03	1.17	1.54	1.10	8.80	3.38	1.07	7
Ž :	/4/74	70104/	1 6	7.5. AB AC	90: <b>.</b>	2.49	2.16	1.72	2,64	4.52	<b>₹</b> .58	~
<b>3</b>	743230	AA400234	0.00	26.36	3.02	2.74	2.82	1.36	2.38	4.05	£.	2
?	000107	#800AL	į	3	;	i	ç					
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Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	IMAGE	Gen Bank Accession	AveBPH	AveTumor	Fold(aveTumoriav Fold(MPM120/A Fold(MPM121/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A \$ of Tumor abph) veBph) veBph) veBph) veBph) veBph) veBph) veBph) veBph)	Fold(MPM120/A (	Fold(MPM121/A	Cold(MPM124/A	Fold(MPM122/A veBPH)	Fold(MPM123/A veBPH)	Fold(NPM126/A veBPH)	# of Tumor Samples
		Number	;	į		. 80	4 37	8.10	2.01	2.27	1.23	7
4	357628	W94121	0.12	0.37	5.0 5.0	2 7 7	5.33	1.83	2.67	2.63	3.85	7
Ŧ :	701766	A292655	7 .	17.71	10°5	2.24	4.24	4.48	1.57	2.77	2.74	7
Ĭ	183407	K82103	3 3	3 6	28.0	1.42	2.73	1.25	1.78	5.99	4.80	7
} }	786730	0780/074	5	737	50 Z	4.17	6.87	1.88	1.67	2.47	1.08	7
1 5	344913	AA036804	2	0.18	2.87	2.12	6.90	1.18	4.61	0.14	2.91	7
7	34680	WOORA	3 6	22.52	2.87	2.25	2.78	1.05	2.96	4.30	3.91	7
3	447700	D18782	. 6	8.21	2.87	2.52	2.84	2.73	4.48	3.12	2.14	~4
<b>1</b>	50530	117010	8	5 5	2.80	1.02	2.38	1.71	3.89	6.14	2.52	2
3	208082	180583	2	12.48	2.86	2.26	6.36	2.48	4.87	0.81	1.18	7
3	400826	4412848	100	0.20	2.85	0.16	1.59	10.45	1.7.1	0.15	3.62	8
3 7	264167	NO CONTRACTOR	8	0.28	2.85	4.52	5.80	2.94	2.25	0.12	2.05	7
452	13340	R27450	122	5.05	2.5	2.28	2.49	1.50	5.90	3.21	2.23	7
157	416889	VARBA32	29.0	1.62	2.92	1.81	5.15	4.77	1.53	1.35	2.83	~
3	271078	N29018	2.08	60.09	2.82	4.05	4.90	2.55	2.04	1.62	2.38	7
455	452320	AA704777	1.07	3.13	2.82	2.22	2.90	2.18	3.07	2.67	4.48	~
85	814798	AA455235	25.12	73.30	2.82	3.83	7.12	1.66	2.07	1.28	1.58	~
157	417908	W80520	5.97	17.42	2.92	2.62	6.08	2.91	4.81	0.62	0.47	7
458	745283	AA625567	5.50	16.00	2.91	2.17	1.35	1.50	2.45	5.86	4.13	~
657	50772	H16803	10.55	30.4	2.80	3.09	5.75	2.12	2.83	1.37	2.15	~
97	51606	H18936	990	1.96	2.88	2.55	3.73	3.99	2.18	2.15	2.70	7
15	204684	H57135	0.83	2.38	2.88	1.27	2.47	2.59	4.88	4.08	2.03	7
462	731002	AA418884	1.1	3.20	2.87	2.09	3.10	4.25	2.58	2.78	2.40	~
\$	124891	R06119	0.85	2.43	2.87	2.68	3.84	3.59	2.24	2.73	2.16	7
\$	428138	AA002007	0.55	1.58	2.87	2.46	3.04	2.60	5.16	2.82	<b>1</b> .11	7
465	45687	H07934	1.59	4.82	2.85	2.11	5.17	3.34	2.83	1.81	2.04	7
8	731264	AA420989	500	0.13	2.84	2.74	1.81	3.99	1.81	2.21	4.61	~
467	46055	H08850	0.74	2.09	2.84	0.41	0.65	0.73	4.40	10.07	0.75	2
468	626385	AA189113	29.81	25	2.63	2.15	2.80	3.18	6.80	1.20	0.70	~
469	276450	N34824	0.47	1.32	2.82	6.04	3.19	1.65	1.41	2.48	2.18	2
470	480435	AA677561	3.97	11.17	2.82	2.79	3.27	2.11	2.34	4.55	1.65	2
471	41825	R54109	0.07	0.21	2.82	0.83	4.13	2.29	2.36	0.14	7.16	7
472	682056	AA258458	1.86	5.21	2.80	4.23	4.71	2.40	2.75	0.67	203	8
Ê	754408	AA436187	0.03	80:0	2.78	5.62	0.31	1.97	2.18	4.95	1.67	~

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	IMAGE Clone ID	Gen Bank Accession	AveBPH	AveTumor	Fold(AveTumor/Av Fold(MPM120/A Fold(MPM124/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A # of Tumor ebph) veBph) veBph) veBph) veBph) veBph) veBph)	Fold(MPM120/A veBPH)	Fold(MPM121/A veBPH)	Fold(MPM124/A veBPH)	Fold(MPM122/A veBPH)	Fold(MPM123/A	Fold(MPM126/A veBPH)	s of Tumor Samples
į		Jegunn.	40 40	40.00	7	2 13	2.47	2.22	3.18	4.00	2.68	74
<b>:</b>		H28213	2. S		5.5 E.C		3.14	4.59	2.82	1.71	2.89	7
6) <del>4</del>	780087	0412/N	8 5	3.45	2.78	3.20	2.91	2.93	0.95	2.36	4.21	8
Ę	406876	D00004	9	184	2.78	68.0	2.80	4.88	2.49	2.42	3.19	2
7 87	731048	A4421278	200	0.05	2.76	0.58	2.17	0.86	4.74	2.38	5. 2.	7
470	812985	AA484800	78.0	241	2.76	4.36	1.66	2.84	3.65	1.52	2.70	~
£	809789	AA454758	0.17	0.48	2.78	1.66	5.83	2.80	3.4	1.72	1.10	7
- 57	289513	N62745	0.02	0.05	2.75	0.51	3.74	2.54	0.51	2.28	6.97	8
5	327239	AA284307	0.50	1.37	2.75	2.08	1.72	3.12	1.74	1.08	6.80	8
3	23588	R38369	1.07	2.98	2.75	2.22	4.10	3.71	2.22	1.48	2.78	8
3	183617	B48813	127	3.49	2.75	1.72	3.42	2.33	2.85	1.79	4.41	8
484	401435	AA150475	0.28	0.78	2.75	0.93	4.33	2.24	4.29	1.74	2.98	7
3 5	236282	H61103	0.12	750	2.75	1,13	0.08	0.98	0.55	8.93	4.84	7
3 5	781342	AA448390	0.08	0.22	2.75	2.79	0.80	1.58	3.17	0.83	7.34	7
<b>3</b>	788525	AA452113	0.61	2.21	2.74	7.49	3.35	0.82	1.98	1.28	1.51	7
3	742082	AA405789	0.13	0.35	2.74	2.57	4.19	4.36	2.31	1.07	1.93	7
8	39821	R53428	12.95	35.29	2.73	2.05	2.54	2.17	3.06	4.23	2.31	7
164	40338	R53072	2.38	6.50	2.73	2.54	4.72	4.92	2.58	0.83	0.77	7
2	194638	R84398	10.30	28,05	2.72	2.21	1.67	1.84	1.82	4.30	4.69	8
\$	149406	H01518	3.58	9.74	2.72	1.80	2.57	3.46	4.97	1.74	1.77	7
å	74223	148412	6.45	17.51	2.71	3.86	6.49	1.22	0.80	1.93	<del></del>	7
485	454798	AA677285	0.71	<b>1</b> .	2.71	2.51	1.34	3.6	1.20	0.70	7.48	7
\$	428338	AA005153	6.3	3.51	2.70	1.67	3.06	2.89	2.48	3.61	2.51	74
787	853505	AA863556	0.52	1.40	2.70	18.1	2.59	3.68	1.98	2.28	3.86	2
80	261204	H98218	0.05	0.14	2.70	4.18	4.07	1.82	2.64	1.18	2.32	64
6	1538971	A017332	1.32	3.57	2.69	1.60	2.33	3.00	5.58	1.99	1.66	7
8	282399	N68399	2.00	5.38	2.69	3.78	4.41	1.27	2.53	1,43	2.71	7
8	876281	AA670286	4.02	10.81	2.69	2.74	3.62	1.27	1.97	4.20	2.33	7
203	194384	H50877	14.85	40.11	2.68	1.36	3.04	2.45	1.92	4.73	2.60	8
g	625824	AA186885	90.0	0.21	2.68	0.63	2.98	7.53	3.18	0.92	0.82	2
Š	810328	AA484143	0.28	0.75	2.67	3.26	2.72	2.12	2.74	2.16	9. 2.	8
Ş	50114	H16743	2.07	5.53	2.87	2.59	6.02	0.45	48.5	0.35	0.70	7
808	743731	AA628357	0.28	0.78	2.87	2.85	1.51	4.67	0.86	4.27	1.88	7
204	452047	AA707118	0.41	1.08	2.67	1.41	6.37	1.25	3.28	1.41	2.29	71
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Table 2-1;T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

		Gen Bank			FOR THE PROPERTY OF THE POST O	EcidimpM120/A	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A	Fold(MPM125/A	# of Tumor
	MAGE	Accession	AveBPH	AveTumor	-old(Ave.) umomy •BPH)	veBPH)	VeBPH)	VeBPH)	veBPH)	veBPH)	V•BPH)	sembres
		Number				ď	200	1.71	1.78	3.89	4.19	
808	381587	AA016980	0.88	2:34	2.85	57.7	70.7		, F.	3.31	2.24	7
809	283142	N51335	4.33	11,49	2.65	1.58	2.50	0.83	8 3		8	8
013	202	W88294	0.58	1.53	2.65	1.63	2.24	3.17	6.41	CF. 6	86. 6	
3	2000	4444080	2	0.83	2.65	4.40	3.73	2.69	2.49	0.20		
Ē ;	20102			0.00	2.85	7.24	3.68	2.09	1.38	<b>7</b> 8.0	0.00	<b>v</b>
912	30008	W/3/300	3 :	5 6		3 03	4.78.	1.15	2.48	<u> </u>	3.00	N :
513	896318	AA598828	2.48	76.0	50.7	\$ <del>*</del>	3.88	3.91	1.86	1.88	2.62	~
614	241787	H91721	0.16	0.42	2.65	1.12	3.48	3.78	2.	3.28	1.73	7
515	1636108	AI015679	2.30	8.08	2.65	2.30	9, 1	2 13	8	2.78	3.37	ч
616	201172	R98487	0.67	1.78	2.65	1.56	<b>4</b> .10	7 7	Į.	4.53	4.38	7
517	228280	H63865	0.70	1.86	2.64	1.33	2.40	<b>t</b> 6	÷ ÷	0.26	8.46	7
618	342164	W83778	<b>0</b> .0	0.10	2.64	0.60	0.26	80.0	• · ·	2	0.63	8
519	49394	H15655	0.51	1.35	2.64	2.46	4,41	2.08	0 5 0 6 0 6	, «	2.30	7
520	85320	171889	6.68	17.60	2.64	1.58	1.93	<b>4</b> .01	28.5	900	7	2
521	115143	186708	5.38	14.12	2.63	3.65	6.57	1.85	C9.1		4.28	. 60
3	31060	PA1004	0.74	1.96	2.63	1.11	1.08	1.83	1.88	3.02		٠,
1 5	202403	07781044	6	25.98	2.63	2.80	4.67	2.27	3.51	1.03	Pr. 1	
3 5	90000	B10004	0.35	0.91	2.63	1.09	4.25	1.50	7.15	0.02	77.0	۷ ،
5	1		3 6	1 92	283	1.79	3.06	2.41	3.54	2.21	2.78	<b>v</b>
9	20715	04742		10.27	2.82	3.22	2.98	1.49	3.17	2.47	240	2
979	71129	1410//	78.6		28.	2.86	2.18	3.18	1.50	2.17	3.98	7
27	49728	H29198	0.70	3 ;	10.2	2.25	2.74	3.72	1.30	2.40	3.25	6
<b>828</b>	295723	N66925	7.87	1.3/	10.7	27.6	3.25	1.52	4.68	2.16	1.28	2
528	447715	AA702781	<b>7</b> .8	38.63	197	6.10	3.75	-	2.53	3.64	3.36	7
S S	898070	AA698779	0.92	2.40	19.2	9. c	2.13	1,	2.01	3.34	3.66	~
531	60540	T40531	1.12	2.81	7.60	2.02	2.5		280	2.10	2.08	2
632	288999	N62728	3.95	10.29	2.80	2.27	3.83	<u>}</u>	<b>4</b> 80	1.96	2.40	7
53	290348	N62293	1.53	3.88	2.60	1.93	9	2		3.45	2.14	8
캶	51865	H23187	0.05	0.12	2.59	2.52	1.63	0	9 6		2.42	8
833	815124	AA481491	1.03	5.02	2.59	1.28	1.68	2.60	n (7)	- ;	3.67	~
853	428296	AA004944	0.60	1.56	2.59	1.85	2.54	2.85	<b>3</b> 0.	2.5.0	5 6	، ۱
63	71807	H19427	0.15	0.40	2.58	2.50	5.88	3.00	0.82	2.24	9 (	4 (
3 5	787481	NEOR47	2	917	2.58	1.74	2.31	0.18	0.37	7.68	3.72	7
2 5		164367	200	9 10	2.68	1.69	5.87	2.69	3.88	0.14	0.82	N
a c		2000	2.5	27.60	2.58	3.63	2.79	2.02	4.23	1.30	1.48	7
3	1647	*20401	2	10 4	2.57	3.34	0.28	0.12	0.68	10.78	0.24	~
Š	3	780/N	2	<b>!</b>			ç					
							16					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

umor	Samples	~	•	•	••	•		•	•	•																									
VA # of T	Sam	٠,٦	~	14	2	N	.,	**	**	14	8	7	7	2	8	2	7	7	7	7	7	7	~	8	2	7	7	7	7	7	7	8	7	8	7
Fold(MPM126	veBPH)	1.95	1.15	1.13	2.23	1.65	1.48	1.17	2.53	3.87	4.51	2.72	3.60	1.59	2.38	1.58	3.89	0.85	2.38	<del>7</del> .	2.32	2.05	1.57	<u>.</u>	1.79	1.27	1.81	1.69	2.48	1.75	2.96	 3	1.05	0.76	±. ₹.
old(MPM123/A	veBPH)	6.79	4.97	1.36	5.51	3.57	3.25	1:1	4.73	5.94	2.81	3.30	3.54	1.48	1.68	5.24	2.52	<b>3</b> 5.5	1.73	1.90	3.20	3.36	1.78	0.86	4.77	3.11	2.62	3.28	3.94	2.69	3.20	1.85	3.01	1.93	1.28
Fold(NPM122/A F	veВРН)	3.37	4.68	3.13	1,45	2.31	0.54	3.32	1.95	1.41	1.04	2.49	2.48	4.20	1.88	1.58	3.11	1.64	2.56	3.58	1.83	3.11	2.23	1.30	4.19	2.90	1.29	4.87	3.23	3.67	4.47	3.20	4.55	2.67	3.78
Alectronic Linear	veBPH)	1.84	1.52	2.87	3.19	2.59	2.09	2.79	3.06	1.28	0.80	0.81	1.99	1.78	3.14	1.05	2.26	1.93	1.80	127	3.26	2.87	96.0	1.90	1.00	4.62	3.63	2.43	2.43	3.08	1.39	2.41	2.81	3.13	222
4/70711111111111111111111111111111111111	rold(MFM1211A	1.51	181	2.47	1.70	3.53	5,19	2.29	8.	1.10	2.89	3.35	2.86	3.69	2.61	4.26	1.48	3.90	3.59	3.50	1.78	1.90	4.71	6.33	1.80	60.0	3.56	1.30	1.68	2.39	1.10	2.83	2.4	4.23	3.42
	Fold(IMPM120/A VeBPH)	88	1 22	84	<u> </u>	2	2.80	89	1.19	1.72	323	2.58	72.0	2.50	3.54	<u>*</u>	1.95	2.23	3.12	2.94	2.69	1.77	3.86	3.52	55.1	2.80	2.26	1.43	1.18	1.32	1.78	3.17	1.01	2.13	2.30
ENGLINE ENGINEERS CALINDRISTA FOLDING	Fold(AveTumor/Av eBPH)	787		7 5	<u> </u>	25.0	5.30 88.00	3 6	35.6	2.45	2.55	2.54	2.54	2 554	2 54	254	2.53	2.53	2.53	2.52	2.51	2.61	2.51	2.60	2.50	2.50	2.50	2.50	248	2.48	2.48	2.48	2.48	2.47	2.47
	AveTumor	6	ac's	8 6	A 5	63.63	, c	· •	 	j 5	2 6	3 6	5. c	4 4 7	15.28	7	08.0	8	2.01	8.38	8.15	1.26	3.50	29.92	6.89	0.27	₹.89	2.74	2.35	8.13	1,83	1.7	1.18	0.83	24.32
	AveBPH	•	ار ای	2 C	) ()	3 5	) () ()	9 6	3 6		7 2	5 6	2 5	2 6	3 20	250	3.91	41.0	0.80	1.74	3.24	05.0	1.39	11.96	2.78	0.11	8	1.10	9	2.47	0.74	0.69	0.48	0.38	9.84
June Bank	_		AA705525	H94471	AA48883	AA868750	H97186	101001	W/4/20	07755W	7440300	707107	1/7704	CARON	TR1038	WAROIA	A A B B 4 7 0 4	AA443140	TB2072	AA025538	AA099593	AA889397	H88329	R25.823	AA676740	AA158030	N35922	W73882	AA882200	AA043552	AA703048	AA489639	R2208B	H17921	N68132
	IMAGE Clone ID		482354	243159	824896	884283	260707	/RLD+L	344018	27,0030	20110	20027	177087	770000	70005	101455	44KRR7K	796723	85870	365738	489722	1468074	252683	36393	460039	590148	272600	345559	1481228	487723	438535	823694	130758	50835	276483
		;	. <b>2</b> 42	£ .	<b>*</b>	5	ž :	3	Š	<b>A</b>	2 2	2	700	200	, K	3 5		-	550	\$ 50	25				3	986	287	288	-		14	572	27.5	57.4	575

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	1	Gen Bank			A land and the second	F-12.8T-1881-15 ENGINEM 2018 ENGINEM 221/8 FORINPM 124/8 FORINPM 122/8 FORINPM 123/8 FORINPM 125/8 # Of Tumor	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A	Fold(MPM125/A	# of Tumor
	Clone ID	Accession	AveBPH	AveTumor	Policia de PH)	veBPH)	veBPH)	veBPH)	VeBPH)	veBPH)	VeBPH)	Samples
953	10100		080	7 0	2.47	3.23	5.03	0.73	2.49	2.35	1.00	- 7
3 5	308483	NORAGE	4	19.63	2.47	2.01	3.49	2.05	4.83	1.49	0.85	7
	785840	20070744	5 5	0 0	2.48	3.20	6.99	1.70	2.59	0.95	0.35	7
8 8	20403	HK244R	0.15	0.37	2.48	1.18	1.24	4.12	5.65	0.95	1.63	7
5 5	2,80570	NATTAR	6 23	45.34	2.46	1.31	3.10	3.00	3.31	2.60	1.48	7
}	A7060	70438	0.82	1.53	2.48	7.96	3.28	2.	0.96	0.82	17.0	7
§ §	25081	770878	14.58	35.84	2.48	1,10	3.69	3.7.	1.95	2.40	1.88	7
\$ \$	100.00	44151017	2 0.7	7.28	2.45	2.15	1.50	2.68	<del>1</del> .8	3.10	4.28	7
3	745208	AAR2RR4	0.53	8	2.45	1.24	2.32	2.05	2.52	3.50	3.07	74
3 8	785890	AA440362	6.28	16.39	2.45	3.01	2.78	1.45	1.12	2.10	4.21	7
8 8	3KB048	WOLKO		148	2.45	3.60	6.39	1.82	1.24	1.17	0.50	7
3 2	487880	180444	5 6	2	2.45	3.25	0.47	1.58	0.77	1.51	7.13	7
3 \$	705407	AA453518	0.13	0.32	7	0,33	0.97	1,47	5.45	0.62	5.82	7
3 8	1480005	AAROOOR	5.05	14.63	2,44	2.08	2.15	1.42	2.58	3.31	3.12	7
9	753162	AA400457	2.39	20.0	2:4	3.25	3.62	0.79	1.82	2.27	280	8
ě	712499	AA278382	2.63	6.41	2,44	1.50	3.82	2.40	3.47	1.58	1.88	7
592	489656	AA099369	8.0	0.10	2.44	4.08	1.37	2.01	1.00	0.24	5.82	~
203	110503	T89996	0.60	1.45	2.43	0.73	1.24	0.76	0.82	5.55	5.38	7
3	197465	H52048	0.35	0.85	2.43	1.53	2.49	3.84	0.91	4.24	1.47	~
505	429729	AA011685	1.61	3.01	2.42	2.40	4.68	1.51	3.19	7.	0.83	~1
8	178187	AA421047	0.03	0.07	2.42	0.89	3.84	2.28	2.85	0.33	4.12	7
202	782585	AA47527	0.48	1.17	2.42	3.45	1.49	5.47	1.78	<del>1</del>	08:0	7
90	811010	AA485385	0.08	0.14	2.42	1.99	2.41	3.45	1.87	0.58	4.13	~•
95	743445	AA608365	3.21	7,78	2.42	2.19	3.40	1.70	1.31	2.63	3.33	7
8	307328	N93470	0.03	0.0	2.42	2.00	3.80	0.67	1.74	2.14	1.15	7
90	297439	W03687	24.42	29.00	2.42	2.70	4.38	66.0	4.78	1.41	0.25	2
602	245320	N53456	1.20	2.89	2.41	4.58	3.25	2.53	2.71	0.66	0.75	7
8	140057	R65963	0.75	1.82	2.41	1.19	2.77	2.60	3.00	1.88	3.13	7
ğ	433294	AA699707	1.60	3,85	2.41	1.23	3.14	1.87	1.24	<b>4</b> .08	2.93	2
808	812227	AA455369	4.47	10.79	2.41	3.67	2.67	2.04	1.25	0.84	0.98	2
808	1560977	AA975010	1.07	2.57	2.41	2.29	2.67	3.01	3.69	1.52	127	7
50	283379	N52767	9	0.10	2.41	4.36	4.19	2.02	0.77	0.87	2.25	~
808	85195	171380	0.02	90.0	2.41	4.76	0.45	0.29	0.45	8.08	0.45	~
809	913310	AA455164	3.54	8.52	2.40	2.76	4.07	Ŧ.	2.17	3.03	0.99	7
						•	18					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

ð	Gen Bank			Enjalana Timodlav Boldinphi120/A Foldinphi121/A Foldinphi124/A Foldinphi122/A Foldinphi123/A Foldinphi126/A # of Tumor	FoldmPM120/A	Fold(MPM121/A	Fold(MPM124/A	Fold(MPN122/A	Fold(MPM123/A	Fold(MPM125/A	# of Tumor
Accession		AveBPH	AveTumor	•BPH)	veBPH)	veBPH)	WBPH)	veBPH)	veBPH)	(H.)	ender se
Number	<b>=</b>	•	;	ç	48.4	2.06	3.15	5.03	<u>18</u> .	0.69	7
H85345	52	<b>3</b> 0.74	48.65	7.40	3 3			1.66	4.37	1.35	7
AA058534	534	0.42	1.02	2.40	2.31	3.10 6.00	<u> </u>	2.47	3.18	4.01	7
AA883314	314	1.81	4.36	2.40	2.09	0.93	2.72	2	3.19	1.14	~
926	AA976714	5.57	13.37	2.40	2.02	2.10	8.6	3	1.47	1.39	2
R93727	.57	1.09	2.61	2.40	4.02	3.00	<b>a</b> (	, ;	7	1.93	7
2	AA420983	3.88	9.20	2.40	5.00	3.67	09.L	70.	1 8 4	2.01	~
Ξ	AA115300	1.00	2.39	2.39	3.92	3.62	1.47	5/.L	0.0	4.48	
ğ	WOZZIA	0.76	1.81	2.30	1.52	1.61	3.50	2.18	90.4	9 9	, ,
Š	A A 37778 27	0.10	0.24	2.39	3.36	2.17	1.55	5.28	0.80	97.	4 6
3	7 5	2 2	87	2.30	3.46	6.52	\$.	1.41	1.37	0.52	7
5	AA418828	47.0	9 5		5.17	3.13	2.98	.0.88	0.70	1.47	7
<b>5</b>	R61796	18.69	4.60	BC.7		333	2.18	3.03	2.42	1.68	~
8	AA005388	1.75	<b>4</b> .18	2.38	60.E	<b>1</b>	99 F	2.49	0.50	0.73	~
35	N35907	11.37	27.10	2.38	10.7	- L	4 4	17	2.81	1.24	2
2	R07891	ন্ত্ৰ	2.83	2.38	3.82	3.35	6	- e	1.43	0.81	~
3	AA458135	12.87	30.59	2.38	1.77	3.33	<b>4</b> 6.7	4.20	4 9	4.16	. 7
3	AA461506	0.03	0.07	2.37	0.67	<b>4</b> .90	0.76	5.01 10.6	: ·	 	۰ ،
ĕ	AA988920	1,50	3.56	2.37	2.49	3.07	2.31	4.6	7.7		۰,
ğ	AA054585	0.20	0.47	2.37	1.64	2.34	3.45	3.39	0.00	2.7	۰,
3	AAR83812	0.78	<del>1</del> 8.	2.37	1.84	3.09	1.87	3.81	06.2		4 (
5 5	BARKER	1.08	2.58	2.37	1.06	1.93	3.22	1.78	4.47	4.74	4 (
f à	4 4 40 40 40	20.5	18.84	2.38	4.09	4.25	2.31	1.16	1.43	6.0	, ,
3	200000000000000000000000000000000000000	18.0	83	2.38	40.	3.55	0.84	0.00	1.70	5.24	N (
3		5 6	78.0	2.38	1.88	3.50	3.02	1.59	1.37	2.79	7
Ţ	Alchon A	2 6	2	2.36	18.	3.27	2.31	2.33	124	3.15	N .
5	78.5		£ .	238	1.65	5.16	4.08	1.43	3.15	1.69	2
ğ	282902	3 3		, c	4.14	3.65	1.91	2.15	0.83	1.43	7
9	AA504783	<u> </u>	9, 0,	, c		4.36	1.89	4.01	1.75	0.50	7
5	N21237	3	2 ;	30.0	30.7	2.28	1.17	0.79	0.75	4.71	7
<u>Ş</u>	H05734	0.57		2. 25 25. 6	e. €	0.28	3.80	0.62	4.07	2.43	7
3	R54105	90.0	170	8 .	6	3.74	1.37	1.91	57.4	0.72	~4
š	W05828	0.81	.08 .08	<b>5</b> . 2. 3	80.2	7.5	330	1.15	3.46	3.76	71
₹	AA417246	<b>3</b>	1.61	<b>5</b> .7	97.7	<u> </u>	7 7	3.24	3.01	2.78	7
3	VA483932	0.60	1.40	2.3 <del>4</del>	1.52	1.71	B (*)	37.	0.12	2.05	~
g	VA598956	0.05	0.12	2.34	3.49	2.31	Z.08	2 6	1 6	79.0	2
2	AA788970	4.63	10.81	2.33	3.43	7.25	0.74	C.0		<b>.</b>	•
						19					

Table 2-1;T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

		Gen Rank		•	1	A A A C A SI C COMPANY COMPANY	old/MPM121/A	OIGINPM124/A F	old(MPM122/A	Section of the Control of Control	old (MPM125/A	# of Tumor
	Class	Accession	AveBPH	AveTumor	Fold(Ave lumonav eBPH)	VeBPH)	veBPH)	veBPH)	veBPH)	veBPH)	VeBPH)	Samples
		Number				,	, 43	0.85	1.86	4.32	1.76	7
£	242037	H83328	0.38	0.89	2.33	3.07	2.13	9 6	333	3 03	1.50	8
64.5	470914	AA032084	13.04	30.41	2.33	2.36	3.53	<b>47</b> .1	£		4.50	2
3	38280	R40462	0.70	1.64	2.33	1.90	0.47	0.87	0.59	5.5	, t	
3 5	9000	00	- 60	88	2.32	3.30	3.08	2.73	2.02	J.S.C	70.1	• •
3	00000	000/11	9 .	9 6	4.0	1.78	3.31	2.05	2.70	3.18	0.92	7
ž	743217	A400128	3 5	D. 4	, ,	3.15	3.16	0.87	2.32	1.70	2.71	7
2	290841	N71982	n Q	60.01	7 .	3	2.10	1,47	3.01	4.67	1.50	8
<b>9</b>	814585	AA480906	ž	13.75	2.32	<u>*</u> ;	2 6	3,48	2.02	1.79	1.58	7
651	240788	H80215	2.74	6.33	25.23	57.1	5.53	9 0	3 79	2.65	3.37	7
652	34869	R44447	2.35	5.45	2.31	1.25	<del>1</del> .34	<b>3</b> (1)	5. T	1.37	3.63	~
653	51283	H18838	90.0	0.14	2.31	4.10	1.12	1.78	6.		80	,
2	810208	A 4 4 6 4 5 7 7	1.10	255	2.30	1.35	0.98	5.47	1.37	3.40	97.1	• •
<b>§ §</b>		200000	200	0 47	230	1.39	2.38	3.59	5.16	0.48	0.63	•
CCD !	10001	AA401508	5 5	. 6	2 6	3.25	2.34	3.04	2.87	1.74	0.77	~
92	745138	AA828698	12.78	28.30	8.5	3 .	330	2.10	5.47	0.55	0.48	~
657	489078	AA057185	8	0.05	2:30	1.80	5.43	2 68	- E	86.0	1.43	7
858	34108	W73781	3.42	7.83	2.29	8. <del>↓</del>	3.1/	5.50	5 6	\$	1.55	64
628	727192	AA283819	12.78	29.26	2.29	2.09	0.89	224	3.41	3 9	1.60	, ,
999	41541	R52789	27	29.6	2.28	2.19	3.57	3.59	1.28	84.	20:-	• •
3	704363	44453487	8	9.86	2.29	1.68	4.15	1.50	4.58	0.88	<b>4</b> 5	٠,
3 8		00000	5	1,0	2.28	2.41	0.86	1.19	0.31	4.68	4.26	7
8	3 5	H32000	5 6	- 6	80.0	1.85	3.18	2.62	2.50	3.07	0.48	7
8	48330	3484	÷ 6	9 6		87.6	0.12	2.08	2.75	5.09	0.47	~
ş	169600	<b>AA425900</b>	3	0.20	07.7	) (1 ) +	3.50	328	1.78	1.88	1.51	~
665	112885	T87139	2.01	5.85	27.78	RC:	9 6	8 6	3.14	0.18	2.05	2
8	362785	AA018683	98	0.13	2.28	\$5.7	B0.0			117	181	7
687	384397	AA708676	0.48	1.05	227	1.47	3.59	4	08:1			
888	26568	R39111	97.0	1.68	227	3.15	0.55	0.70	7.84	3 5	9 9	
880	100228	1005811	65	1.27	227	1.69	3.64	3.07	1.83	3	<b>c</b> 0.1	4 (
22	44070	BARTON	50.0	0.07	2.27	1.48	2.18	0.33	1.19	4.15	4.27	<b>7</b>
3 6		2000	,		2.97	0.74	4.79	<del>1</del> .	1.55	3.13	<u>a</u>	7
6	50110	117911	5 6	3 3	70.6	3.52	25.2	3.03	2.05	0.61	1.8.1	2
7/0	346557	N. A.S.	70.0	5 6			3 20	1.52	2.48	1.77	1.51	7
Ę	624785	AA181995	<b>3</b>	70.8	7 7 6		93.6	ă	2.27	1.42	1.87	7
974	626716	AA191548	17.62	38.85	97.7	8 8	9 6	17.0	2.68	0.71	0.53	7
875	785760	<b>AA449686</b>	0.05	<b>5</b>	278	90.5	B (*)	- 97	1 02	090	1.62	2
676	345145	W72231	3.57	8.07	2.28	3.73	3.52	61.7	78.7			,
677	46820	H10036	15.87	35.82	2.28	96.0	1.62	1.52	1.48	3.02	<u> </u>	•
						•	20					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

		S S S S S S S S S S S S S S S S S S S			# of Tumor   Sold(MPM120)A Fold(MPM121/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A \$ of Tumor   Samples   Samples	Fold(MPM120/A	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A	Fold(MPM126/A vebph)	# of Tumor Samples
	IMAGE	Accession	AveBPH	<b>AveTumor</b>	•BPH)	veBPH)	veBPH)	Vebru			1 07	2
_	Cione ID	Number					120	2.33	0.21	9.63 6.63	0.	
į	044823	7082878	0.05	0.11	2.25	0.8/		3.18	1.73	1.38	3.78	4 (
8	100110	Liser K4	0.19	4.0	2.25	1.32	11.7		1.38	1.62	2.28	N
2/9	41.04	Lange L		5.67	2.24	3.20	3.06	70.1	75 6	0.37	0.36	~
8	823691	20/884AA	3 7	7.92	2.23	2.21	5.87	9 6	3.17	7.7	1.76	2
	754127	AA478627	<b>5</b> 5	=======================================	2.23	1.70	<del>1</del> .3	g Ç	55.6	0.72	3.	64
	1461609	AASSJOU	8 6	0.70	2.22	1.86	4.17	7 - 7	1	1.67	1.72	7
683	42088	H05140	4.5	48.25	2.21	3.16	3.03	<b>1</b>	348	0.79	1.35	7
\$	232012	H/2643	3 5	0.25	2.21	1.97	3.06	8C.2		1.72	0.28	7
8	135789	K33355	. 0	073	2.21	3.92	3.22	J. 16	2.25	1.21	3.10	71
<b>8</b>	905097	AA404148	2 6	0.13	2.21	2.46	1.00	5.42	2.18	2.77	3.05	2
687	0/28*	AA111080		3.77	2.20	1.65	3.01	6.3	517	0.91	3.26	~
3	250707	Accessor	56	10.05	2.20	1.61	1.24	1.02	2.10	1.98	08:0	2
3	2000	10724	98 0	2.18	2.20	1.77	3,15	- 6,	908	1.59	3.27	7
8	17557	484//L	1 49	3,27	2.20	2.38	1.42	8 + 1 · 0	2 S	0.20	1.07	7
68	2002	000000	0.13	0.29	2.20	1.26	3.11	0,7	136	3.30	3.35	7
8	78254	AAAAA	2 6	90	2.20	1.94	2.03	BL.F	3 5	1.12	1.09	7
<b>3</b>	813041	20000	2 6	8.47	2.19	2.26	3.85	3.24	6.0	0.20	5.50	7
ĝ	126588	70690X	9 6	11.0	2.19	1.87	<u>r.</u>	3.63	27.6	3.28	0.88	7
685	345553	W73858	3 6	7 40	2.19	1.10	1.74	2.61	2 t. c	1 83	6.60	8
8	51947	H24313	60.7	120	2.10	0.84	0.11	3.51	2 6	4.87	0.81	7
697	<b>4</b> 0010	R54036	2 6	3	2.18	1.56	3.14	2.20	0.93	135	2.78	7
88	383188	AA074224	87.0 90.0	5 5	2.18	3.98	1.50	0.27	3.22	0.78	1.83	~
689	73531	155639	60.0		2.18	0.48	0.82	6.02	<b>t</b> 0.0	1.24	1.53	2
ğ	768020	AA18748	3.03	3 5	2.18	2.11	3.23	3.65	8Z.T	7 6	0.77	~
5	33860	R44840	G. 13	2.5	2.17	4.03	4.62	2.45	1.01	77.0	2.79	7
ğ	46166	0/0804			2.17	3.17	3.06	0.65	7.40	70 6	1.08	2
703	283191	N51382	7/7	2. S. A.	2.17	1,43	3.18	0.73	2.68	<b>.</b> 5	4.23	7
ş	647985	AA207105	• F	12.38	2.17	1.28	0.93	1.27	0.1.0	1.00	1.39	~
55	491001	UL/981W		234	2.16	1.67	1.35	3.92	0.4.0 0.00	9 E	0.68	7
8	1481181	AABGBU3B		7	2.18	1.68.	4.53	1.35	3.60	27.0	1.23	7
707	757503	AA420300			2.18	4.65	3.70	1.70	08.0		0.82	7
<b>28</b>	280375	X47113		5.4	2.18	1.68	3.74	1.81	3.80	9 6	80 6	7
709	363916	AA021247		B 04	2.15	1.34	3.07	1.77	1.30	2.30	3.20	8
710	121729	T98001	_	2.3	2 45	1.58	0.79	0.98	5.28	90. 1.0	, ,	
7	685801	AA262080	1.29	7.7	<u>:</u>		27					
							i					

Table 2-1:T3NO vs. BPH; Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

		Gen Bank			FINAL TO THE POLITION FOLD MIND MISSIA FOLD MIND MISSIA FOLD MAN MISSIA FOLD MIND MIND MISSIA FOLD MIND MIND MISSIA FOLD MIND MIND MISSIA FOLD	EnidimpM120/A	Fold(MPM121/A	Fold(MPN124/A	Fold(MPM122/A	Fold(MPM123/A F	old(WPM125/A	# of Tumor
	MAGE Clone ID	Accession	AveBPH	AveTumor	FOIG(AVE LUMBOILAY	VeBPH)	veBPH)	VeBPH)	veBPH)	veBPH)	VeBPH)	SeidEs
		Number		!		,	1.36	1.33	3.92	1.95	3.18	
712	207370	H58834	0.13	0.27	2.15	• :	3 6	,	227	1.68	0.96	2
713	202795	H53864	2.15	<b>4</b> .60	2.14	1.36	5.0	8 .	97 +	10.0	2.0	7
714	826138	AA521337	0.01	0.02	2.14	4.28	<b>4</b>	17.4	2 6	1.30	1.13	8
715	322481	W15305	2.11	<b>4</b> .50	2.13	3.57	3.32	2.70		1.16	4.1	~
718	168232	H45711	0.61	1.31	2.13	3.34	<u> </u>	t :	3,4	7	0.47	2
717	197525	H52001	2.60	5.54	2.13	2.40	3.78	3.35	£ 7	103	0.69	8
718	460279	AA677629	9970	<del>*</del> .	2.13	2.50	3.41	1.76	14.5	10.	5	
719	24544	N53482	0.63	1.12	2.13	1.10	2.23	1.40	3.87	5 5	88.	i <b>č</b> v
2	83029	T67807	1.42	3.02	2.13	0.89	3.50	3.33	2.60	0.58	3 2	• ~
ž	276871	N30428	0.05	0.11	2.13	6.86	0.19	1.08	0.19	r.,	, ,	
<u> </u>	11060	1007001	0 12	0.25	2.13	0.40	¥:4	0.31	<b>Z</b>	1.68	80 7	• •
1 1	BD 7 6 6	ine /ow	<u>.</u> :	7 87	2.13	1.47	3.12	3.23	1.23	1.67	2.04	7
2	CC7 H.Z	2000N	2	5.7	; c	1.35	3.02	1.62	1.40	4.42	<b>3</b> .	7
724	243/00	N45083	<u> </u>	\$ 5	; ;	9 - 6	3.00	0.85	1.26	1.60	2.84	~
22	285040	N63375	S .	12.63	2 6	2 5	3.05	1.47	4.55	0.82	1.02	7
128	73252	156013	<del>2</del> .	7.62	7.7	07:1	1 15	3.08	0.82	0.92	5.71	7
12	743485	AAB09385	2.	3.78	11.7	0.35 4.55		2.17	1.16	1.40	1.27	~
728	154485	R54850	4.31	<b>9</b> .08	2.11	3.15	900	9	2.09	0.88	2.01	7
22	814780	AAA55227	0.13	0.27	2.10	3.08	D	2.4	1.85	1.43	3.49	7
ξ.	322005	W37418	0.91	1.80	2.10	1.29	21.1		3 4	1 17	1.68	8
151	416945	W87710	3.08	6.46	2.10	2.	3.31	3.18		3.20	1.03	7
732	624429	AA182680	6.29	13.17	2.10	1.20	3,14	1.32	6.00	93	7.7	•
2	18181	AAB09688	90.0	0.18	2.09	0.17	1.08	0.74	4.67	8. 8	9 6	, ,
2	14517R	W72283	0.18	0.37	2.09	1.24	1.24	1.45	4.29	4.03	0.20	۰ ،
5 5	878A57	44870200	87	3.08	2.00	1.94	3.20	1.64	3.65	0.89	= ;	7
3 5	36661	A A 1788/10	67.3	8	2.08	1.37	3.02	3.06	1.55	1.40	<b>3</b>	~ •
3	316473	2007	7	60.0	2.08	1.48	3.27	3.55	0.87	1.38	1.95	7
3	277900	2000	5	0.13	2.08	2.84	0.16	0.58	3.12	4.83	78.0	7
9	0	C704/N	3 :		906	2 0 7	3.92	0.82	3.60	1.36	7.	7
238	33/204	W83585	- F	07.7	8 6	80	3.85	1.82	0.25	0.28	5.19	~
740		K32640	3	9 1	3 2	8	4.34	1.69	1.59	3.62	4.14	7
Z	462857	AA682425	Ŗį	2.46	2.04	0.87		71.0	0.83	1.18	3.40	7
742	292770	N63646	0.07	0.15	70.0	[ <del>4</del> ,4]	6.33	90.	5	0.32	0.80	7
743	185635	R69317	870	0.63	2.04	3.70	00.0		9	190	0.50	7
Ŧ	663673	AA101299	0.58	1.18	2.03	2.13	3.21	3.13	90.7		183	. ~
745	705522	AA454218	9.0	0.10	2.03	99'0	0.20	0.98	3.23	¥.0	3	
							22					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	1	Gen Bank			Fold(AveTumor/Av Fold(MPM120/A Fold(MPM121/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A	old(MPM120/A	Fold(MPM121/A	Fold(MPM124/A veBPH)	Fold(MPM122/A veBPH)	Fold(MPM123/A F	cold(MPM126/A veBPH)	s of Tumor Samples
	MAGE		AveBPH	Avelumor	eBPH)	VeBPH)		•	;	3.33	1,27	4
		Number		į	6	1.37	1 12	3.98	2 ;	970	4.35	2
748	42035	R59068	0.18	0.38	S 6	4.97	1.05	0.57	<b>5</b> 8.0	4 27	3.89	7
747	280412	N47208	0.02	0.09	5.03	5	0.21	3.77	1.83	. ·	3.88	7
748	700302	AA283893	Š	0.09	203	30.5	0.81	1.78	0.58	8.	700	7
2,5	40082	R52530	0.33	99.0	2.02	0.60	3.11	1.86	3.12	1.33	, e	. 2
750	1486109	1	112.27	228.02	2.01	60:1	90 6	1.36	1.47	2.10		
2 1	445008		0.27	0.55	2.01	3 1	2 2	1.08	3.17	3.58	177	4 (
2 5	2000	44773177	2.08	4.18	2.01	0.73	5 7	3,05	3.01	1,55	300	4 (
797		4 4 8 2 5 2 0 7	10.80	21.69	2.01	1.7	. Ca	3.51	1,34	1.38	0.77	<b>N</b> (
2		M03447	0.65	1.30	2.01	<u>7.</u> 2.	3.53	1.12	4.05	0.90	4.37	7 (
*			800	0.16	2.01	1.48	0.12	1 60	0.61	2.31	3.41	N (
755			3	0.27	2.01	0.92	3.66		4.80	4.27	0.70	7
758			<u>.</u>	£ 43	1.99	0.61	0.28	1.5.1		1.74	0.87	м
757		_	2.74	÷ ;	1.89	1.28	3.38	1.63	3 6	3.70	0.67	7
758	278498		2.33	3	80	0.56	4.06	1.99	78.7	7	0.49	84
759	32962	R43910	0.33	0.65	96.	1 68	3.65	1.14	3.73	5.1	0.49	2
780		AA670411	13.20	26.15	0A.F	3 7	3.00	1.85	3.68	1.0/	5 6	2
781		_	9.88	19.43	1.97	į ;	44.6	0.45	3.67	<del>5</del> .	25.0	. ~
, E	_		1.76	3.45	1.98	<del>-</del> 57	414	1.49	0.91	4.0	77.	
3 5	•		2.57	5.02	1.95	3.53	· •	1.08	1.57	0.60	3.40	4 (
3 ;			0.13	0.25	1.95	1.25	40.0	1.32	3.38	1.28	1.18 8	N (
\$			65	6.82	1.85	0.96	3.01		1.89	1.68	<u>3</u>	<b>N</b>
8				68.0	1.85	3.04	3.06	9 10	0.63	0.15	4.47	7
8	•	-			35	0.15	3.60	2.65	3 5	. 980	0.07	7
787	7 376043	•	0.01	2.0	- <del>-</del>	2.30	1.24	3.70	3.77	3.22	3.15	7
768	51373			07.0	5 5	15	0.87	1.45	C. 1		1.6	7
769	261363	AA017708		10.65	8 6	0.78	1.30	1.49	1.78	6. 6	99.6	8
130	0 451121	•	0.59	51.1 50.0	<u> </u>	1.31	1.88	0.78	0.75	300	4.76	2
17	1 769833			88.0	÷ ÷	0.80	0.16	1.67	0.16		3.01	8
77	2 782828			0.12	78.1	4 92	0.83	0.83	0.96	A (C)	88.0	7
773	3 841281	1 AA487203		2.93	78.1	. 6	3.63	1.18	3.76	0.32	2.53	~
11.	4 610883		2.84	5.45	1.82	. e	3.09	0.62	0.89	1.23	5.45 0.00	. ^
775		0 AA453783	2.20	4.23	1.92	0.40	0.39	4.59	5.15	0.39	P . C	
17.		8 AA485351	0.03	0.05	1.82	3	0.65	1.41	3.58	0.92	3.71	• ເ
E			5.91	11.33	1.82	7 .	3.20	1.12	3.17	0.78	<u> </u>	٠, ١
77.8			6.73	1.38	1.91	‡. 7 7	97.0	95.0	1.35	3.58	80°C	•
7.		_	900	0.10	-1.80	9.5						
:							23					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

		Geo Beck			Section Commensation Folding M121/A Folding M124/A Folding M122/A Folding M122/A Folding Section Commens	Folding M120/A	Fold/MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A F	old(MPM126/A	# of Tumor
	MAGE	Accession	AveBPH	AveTumor	Fold(AVe Lumon AV	VeBPH)	veBPH)	veBPH)	veBPH)	veBPH)	(T)	
		Number			•		87.0	2.48	3.11	3.59	1.14	
780	208570	H63223	0.59	1.10	1.89	0.51	9 .	2 7	0.87	\$.	3.49	7
781	131887	R25166	2.12	3.98	1,88	3.19	1.11			07.0	1.31	7
787	341978	W81361	128	2.41	1.88	3.07	3.06	1.81	1.0	0 0	0.92	8
	767797	AA682203	1.49	2.80	1.88	3.10	0.77	2.34	3.24	9 9	92.6	~
3		2077	8	K A7	1.87	0.83	69.0	<u>.</u> 2	0.50	0.1.	9 9	۰, ۱
Ž	0000	AA456818	3 8	5 5		0.98	3.01	1.39	4.56	0.80	<b>9</b>	4 (
785	33808	R44538	7	34.4	9.	90.0	3.80	1.51	1.20	0.08	3.55	74
78	25	R4154	0.12	0.23	1.87	C 10.0	0 6	18	3.55	0.24	0.23	7
787	743189	AA401434	<b>5</b> 00	0.08	1.88	2.05	3.30	28	4	1.38	1.8	7
788	247194	N54036	0.38	0.68	1.88	60.1	3.23	2 5	3.07	1.82	0.79	7
789	183771	14448	<u>2</u> .	3.58	1.88	123	٠ د د د	1 6	171	1.48	0.13	7
790	491764	AA150502	0.08	0.14	1.86	3.80	3.08	03:1		0.42	0.42	8
791	782594	AA447531	0.02	<u>8</u>	1.85	3.26	3.78	£1.5		55.	0.39	7
792	471859	AA035144	0.03	0.05	1.85	3.10	5.05	76.0	· ·	0.0	3.28	2
783	278712	N34895	0.13	0.23	1.85	0.30	0.38	5.27	1.73	5 6	0.59	7
2 6	77177	1001	53	2.83	1.85	2.02	3.26	0.79	3.22	95	0.42	
į	261774	HOROKA	2.71	200	2.	1.16	3.48	1.61	00. E	85.0	, e	. ~
2 2	19001	44487808	11.79	21.75	<u>z</u> .	3.03	3.73	0.71	2.01	0.60	9 6	. ~
2 ;	2000	2442444	8	18	2	99:4	0.80	0.64	0.68		90.4	، ۱
	20706	44104070	3 6	3	1.83	1.41	3.32	1.19	3.24	1.13	80.0	<b>4</b> (
2	11011	708001			1.82	1.28	3.14	1.17	<b>1</b> .64	3.17	0.53	<b>,</b> (
2	22020	79/884497	3 3	5 6	£ 5	1.57	0.53	0.62	0.77	4.18	3.24	7 (
8	18083	K15784	5 6	3 5		3.07	0.53	1.15	0.95	69.0	3.60	7
60	358531	W26134	27.7	8 6	9 6	18.0	0.47	3.08	3.10	0.42	2.00	7
807	3233	R42823	7.	0.Z4	9	, q	0.57	900	1.03	3.31	¥.3	~
803	417688	W99074	0.20	0.32	08.1	2 6	7	0.73	1.30	0.25	4.72	7
<b>8</b>	1435339	AA857748	9	0.07	1.79	0.22	76.4	3.38	1.30	0.20	0.43	7
803	128973	R10301	0.02	0.09	1.75	36.0	į (	3.73	3.19	0.89	98.0	7
808	361323	AA017544	2.04	3.51	1.76	<b>\$</b> (	3 5		34	0.20	4.04	7
507	586440	AA148882	0.05	0.09	1.75	0.31	0.70	700		0.90	3.53	2
808	857196	AA773139	0.40	0.79	1.74	1.59	3.02	<b>t</b> . 6	33	0.31	0.31	7
808	500958	AA130964	0.03	90.0	1.71	6.02	rs.0	3.02	7.0	3.11	3.35	7
510	741891	AA401972	5.81	<b>9</b> .	1.69	1.65	69.0	0.62	9.75	0.57	0.18	7
218	610027	AA455275	0.0	0.07	1.69	4.25	0.80	<b>3</b> .00	0 6 6	0.23	1.05	~
812	811609	AA455020	0.03	0.05	1.65	5.05	0.33	120	90.5	0.33	75	7
813	361807	W92431	0.21	0.35	1.63	0.45	3.01	1.71	<b>8</b> /.0	6.5	; ;	)
							24					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	TAN	Gen Benk	į		Fold(AVeTumoriAv Fold(MPM120/A Fold(MPM121/A Fold(MPM124/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A F of 1 umor	r Fold(MPM120/A	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A F veBPH)	old(MPM125/A veBPH)	Samples
		Accession	AveBPH	Avelumor	eBPH)	veBPH)	Vebru				•	c
		Number		,	4	0.95	0.10	0.51	3.09	3.13	S .	4 (
\$14	785538	AA459651	0.0	0.16	6:		890	0.38	3.52	990	3.79	7
815	260168	N32071	0.02	0.02	1.59	i.c.)	3.5	0.43	3.84	0.43	0.43	7
818	609935	AA189154	0.05	9.0	1.59	0.28	2 4	2	0.45	0.16	1.52	8
817	47187	H1040	90.0	0.10	1.57	0.21	3.30		200	0.18	Q.81	7
818	810948	AA459383	9.05	0.08	1.57	1.00	3.48		597	3.05	72.0	7
3.0	490556	AA100696	60	0.07	1.56	3.03	0.23	<u>.</u> .	0.20	0.18	7.0	7
5	789679	AA425826	0.11	0.17	<b>3.</b> 1	4.53	0.39	30.0	1.51	0.22	0.22	7
2	454083	AA676996	0.05	0.07	1,47	3.29	3.40	0.89	0.81	0.31	0.67	7
2	52071	H24350	0.27	0.39	1.47	3.02	3.31	3.30	700	0.33	0.33	7
823	511088	AA100298	0.03	9.	1.35	0.33	0.40 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5	4.50	0.23	0.40	3.07	~
124	742818	AA400482	0.24	0.32	1.33	3.06	0.03 0.03	1.14	2.61	0.17	0.17	
828	504464	AA152178	90.0	1.88	32.02	186.97	10.2	; ;	0.14	155.48	2.32	-
828	306420	N92899	70.0	1.95	26.82	2.37	G .	137 15	0.22	0.14	0.32	-
27	452374	AA700876	3.59	80.73	22.47	0.60	1.30	2.48	0.75	91.20	0.14	-
828	384015	AA702640	0.32	5.24	16.48	2.18	2.14	200	2.14	46.98	2.25	-
828	48841	H10344	0.07	0.64	8.77	0.83	0.14	1 78	41.0	40.15	1.38	-
2	156322	R72818	90.0	0.81	7.60	1.37	0.78		7	0.58	33.43	-
5	81413	T80161	0.15	0.87	6.43	0.62	0.83	9 <del>7</del> - 7	2	2.90	2.23	_
5	745232	AA626152	0.33	1.89	<b>6</b> .03	2.98	2.15	Z4.08	3. 4	0.25	31.39	-
2	428210	AA007276	0.27	1.55	5.77	0.80	0.45	0.30	50.5	69.0	0.65	-
2	50354	H17804	0.28	1.46	5.61	20.0	1.18	0.0	2.5	1.70	27.99	-
	364583	AA022849	0.30	1.05	5.42	0.52	68.0	0.13	1 49	1.10	27.41	-
2	488422	AA044882	900	0.27	5.40	1.15	0.80	0.03	• • •	1.89	24.70	-
2	50799	H17630	0.30	1.59	5.38	0.82	1.21	2:14 	5 5	24.08	1.74	_
8	302180	N78989	0.22	1.18	5.35	2.39	0.79	1.0. 4. ac	92.0	8.	0.26	-
839	128083	R09747	9.0	0.20	5.12	1.88	90.L	26.78	1.46	0.14	0.14	-
3	26933	R36969	0.07	0.37	4.96	96:0	0.62	25.48	0.07	0.08	0.43	-
2	229666	H68542	0.11	0.53	4.77	2.48	80:0 •	22.85	1.24	1.38	0.62	-
72	744627	AA621310	0.82	3.87	4.71	0.95	1.14	33	2 09	22.55	0.40	-
843	884462	AA629707	0.15	0.68	4.42	0.13	1.05	1.35	20.54	0.78	2.08	-
3	361291	AA017468	1.00	4.41	4.41	0.57	C1.1	. 8	0.42	1.17	0.09	-
<b>25</b>	301735	N90882	0.11	0.49	4.33	19.43	50.5	23.85	0.22	0.21	0.41	-
3	753078	AA436479	2.41	10.37	4.30	0.47	6.6	1.32	1.01	69:0	2.52	-
2	510381	AA055585	0.32	1.35	4.23	19.31	16.0	<u> </u>				
							25					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

		Gen Benk			THE STATE OF THE PROBLEMS OF FRICING PROBLEMS AND THE POLICE POLICE FOR THE PROBLEMS OF THE PR	Average 120/A	Fold(MPM121/A	Fold(MPN124/A	Fold(MPM122/A	Fold(MPM123/A	Fold(MPM126/A	# of Tumor
	MAGE Clone ID	Accession	AveBPH	AveTumor	Fold(AVe) umorray eBPH)	veBPH)	veBPH)	VeBPH)	veBPH)	veBPH)	v•BPH)	Samples
		Number			į	č	174	18.40	0.98	2.00	1.12	-
878	448401	AA777538	0.45	<u>1</u> .	4.21	<b>S</b>		2 6	,	10.01	50.0	-
2	504040	AA150819	0.17	0.70	4.16	1.19	1.74	0.82	<b>3</b> :	2		•
95	122298	T99158	0.42	1.7	4.08	1.32	2.06	1.18	1.28	<b>7</b> .	90.01	- ,
1	281442	******	0.39	558	4.03	1.32	1.09	0.51	0.38	20.03	98.0	_
5 5		44034044		60	3.97	1.74	<u>2</u> .	0.57	0.0	17.96	1.89	<del>-</del>
3 5		2000	* * C	2 28	3.82	2.13	2.98	1.30	1.23	14.18	1.67	-
3	7767	107957	9 5	3 7	. E	1 08	1.29	15.59	1.72	2.23	0.81	-
ž,	117470	AA480873	? ;	3. 5	27.6	137	0.95	2.14	0.67	14.52	2.78	-
33	43060	R61868	7.0	2 .	*	04.0	68.0	1.22	1.92	1.22	15.62	-
<b>2</b>	347183	W80666	9.48		0.50 8 8 6	2.08	1.14	1.62	10.80	2.71	2.58	-
Š	32381	K43462	S (	0.00	P	27.0	0.21	1.36	0.63	1.85	16.16	-
85 85	882778	AA158211	3	21.0	D 17	6 6	86.0	1.22	14.84	1.32	2.08	-
839	454908	AA877403	0.60	2.05	3.45	0.77	6.50	9	15.08	0.51	0.39	-
8	340641	W56760	0.27	0.01	3.41	7.57	77.	9. C	800	1.03	0.13	-
8	897722	AA596983	0.08	0.28	3.40	16.70	U.81	0.70	6 6	23	0.05	-
883	866882	AA879352	0.18	20.	3.38	17.69	1.01	0.60	0.60	70	÷ ;	
200	130664	R22579	99.0	277	3.35	1.17	0.97	14.33	121	A.	<u>4</u> 5	٠,
88	1806080	AA995508	0.47	1.58	3.34	1.80	1.58	1.72	1.32	11.93	3 :	- •
885	768260	AA424950	0.82	2.69	3.28	1.66	12.82	2.38	0.89	1.08	23 C	- ,
Ş	46830	H10028	0.28	0.92	3.27	1.16	0.99	0.74	<b>2</b> .	13.18		<b>,</b> ,
} }	1015785	A A R 2 0 1 1 7	10.22	33.26	3.25	0.55	0.26	1.17	15.74	1.49	0.32 23	-
3	40103K	AAAADORB	12.0	0.87	3.24	1.63	0.50	2.32	1.38	2.40	11.24	-
3 \$	104515	BAA242	0.68	1.87	3.24	6.47	2.02	2.78	2.04	2.78	2.43	-
3 6	308413	100000	0 0	500	3.21	1.60	0.62	1.68	0.62	12.88	1.87	-
	2000	000000	27.0	8 8	3.20	1.31	1.70	0.76	2.24	10.28	2.91	-
3 2	PO0081	44455107	900	0.28		0.83	0.72	0.87	<b>7</b> 6.0	0.35	16.41	-
<b>.</b>	27,175	877778	180	6.71	3.5	2.63	6.05	2.60	2.45	2.82	2.41	_
	07.770	NA2251	0.10	0.31	3.14	1.79	0.93	12.98	2.84	0.10	0.23	-
	K424R	74673	<b>1</b>	10	£ 5	1.22	1.79	1.32	1.82	11.65	1.01	<b>-</b> -
	50163	44147714	0.03	0.10	3.13	0.62	0.57	11.65	2.53	1.55	1.85	-
È	81051	104747	0.33	10	3.10	1.50	12.98	1.31	1.08	0.77	8.	-
. K	420175	A & 0.048.24	0.19	0.58	3.08	1.55	6.34	2.94	2.52	2,48	2.84	
878	3448	R44164	0.48	14.1	3.05	0.95	11.69	0.77	1.45	1.05	2.37	-
8	280217	14724	6.29	19.12	3,04	1.96	1.84	0.85	0.83	11.86	1.12	-
2	360838	AA017141	7.0	1.01	3.00	2.60	1.86	2.57	99.1	1.18	8.11	-
							26					

Table 2-1;T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	MAGE	Gen Bank	AveBPH	AveTumor	Fold(ANDM123/A FOLD(MPM121/A FOLD(MPM124/A FOLD(MPM122/A FOLD(MPM123/A FOLD(MPM125/A # Of Tumor 	Fold(MPM120/A	Fold(MPM121/A   veBPH)	Cold(MPM124/A	Fold(MPM122/A I	Fold(MPM123/A F veBPH)	Fold(MPM125/A VeBPH)	# of Tumor Samples
	Clone ID	Number						12 90	1.25	1.05	0.81	-
CRA	730946	AA416543	1.1	3.32	2.98	0.90	08:0	12.67	7.0	1.21	0.61	-
	705207	AA454155	2.01	5.93	2.85	0.60	0.67	5.0	7.41	1.89	1.65	<b>-</b> -
3	400727	B95773	0.39	1.13	2.91	2.12	2.71	1.08	7 7	7.87	1.73	-
	(C788)	C//CBN	90	5	2.91	1.15	2.19	2.85	99.1	2.58	2,12	-
Ş	40,000	AA760021	3 6	8 7.4	2.80	2.81	2.85	<del>1</del> 8.	5.15	2 6	40.2R	-
<b>8</b>	103943	AA279145	5 6	: :	98	1.39	<u>추</u>	1.32	1.43	9.5		
887	781027	AA448031	0.63	8 6		0.52	ġ	0.52	2.66	2.19	40.95 40.05	• •
<b>8</b>	625516	AA187595	0.02	90.0	89.7	7. 6	0.45	1.22	10.05	7.	2.83	-
880	838478	AA457517	<b>4</b> .06	11.68	2.88	1.28	7.47	2.29	1,27	1.75	1.91	-
980	48367	H08959	14.1	¥.08	2.88	75.5		2.53	1.74	7.18	1.78	-
891	213753	H72322	0.28	0.73	2.87	1.57	0.40		1.73	9.18	1.31	
208	128449	R06708	0.22	0.63	2.87	0.97	C7:7	- 6	1.18	2.07	6.38	<b>-</b> -
	204243	V95007	0.60	1.70	2.88	1.99	2.55	08.7	2 5	1.03	1.18	-
2	36447	N22230	122	3.49	2.86	98.0	0.81	12.24	3 .	7.76	1.48	-
	11107	000000	1 6	0.41	2.85	2.05	2.36	1.80	2.13	7	1.98	-
0 9	COZRAL	CROCKY	22.0	2.10	2.84	2.50	2.20	2.20	707	2 2	7.65	-
9	01810	H22303			2.83	1.63	1.87	1.92	<u>z</u>	70.7		-
687	111370	T85173	BQ"	t :	3 2	1.76	2.11	0.92	7.08	2.69	7:30 10:0	• •
868	376435	AA039713	1.18	<b>5</b>	3 8	Ç	2.13	1.85	1.95	7.78	7.07	- ,
888	24676	R36809	0.40	1.13	18.2	2 7	28.0	10.42	1.72	1.24	1.21	- •
8	127831	R08938	99.0	1.85	2.79		040	0.82	2.18	75.0	11.87	-
901	77915	T61323	8.10	22.58	2.79	0.1		2.59	5.69	2.57	1.69	<del>-</del> -
905	327481	W20438	0.76	2.12	2.78	2.05	6.03	1.45	1.03	2.33	1.61	-
903	40263	R55261	0.15	0.41	2.78	1.94	87.8	 	1.86	2.95	5.61	-
8	826109	AA521327	0.40	1:1	2.78	1.65	/A'.	10.0	1.30	1.08	1.28	-
808	113160	T83842	99.0	1.83	2.77	55	56.		1.69	10.48	0.03	-
80	288768	N62498	0.35	0.96	2.77	9.19	0.1		8.37	1.28	1.29	<del>-</del>
200	289938	N58338	0.57	1.58	2.78	1.73	11.7	o		0.69	1.02	-
808	770868	AA434487	0.09	0.28	2.75	12.49	0.40	4 C	17	1.4	1.48	-
000	197778	R63515	<del>1.</del>	3.91	2.74	0.94	97.1	9.75	1.37	1.26	1.62	-
010	140831	R67218	9.8	2.57	2.73	1.07			930	1.90	1.87	-
5	61749	H24320	2.85	7.75	2.72	1.73	1.78	20.7	10.18	0.79	0.90	-
912	280977	N50864	6.	2.74	2.71	1.62	84.1 66.1		7	8.	1.16	-
913	41835	R54212	0.33	0.90	2.70	8.20	1.63	6.10	85.89	1.45	0.60	-
914	951125	AA620558	15.08	40.69	2.70	2.31	2.72	50.6	7.45	2.53	1.28	-
915	51011	H19242	4	10.78	2.68	1.31	0e.r	70.7	?			
							27					

Table 2-1;T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

		Gen Renk			Fold MPM121/A Fold MPM122/A Fold (MPM122/A Fold (MPM123/A Fold (MPM126/A & of Tumor	Alocaldonate	A)14/MPM121/A	Fold(MPN124/A 7	Old (MPM122/A	old(MPM123/A F	old(MPM126/A	# of Tumor
	Gone ID		AVBPH	AveTumor	Fold(Averumon/Av	veBPH)	veBPH)	VeBPH)	veBPH)	veBPH)	VeBPH)	Semples
		_	;	;	ě	1 67	2.81	2.	1.80	5.50	2.76	-
916	44303	H06380	1.99	5.32	2,68	76.7	5 5	320	92	1.88	2.19	-
917	266838	W01211	0.61	1.63	2.68	1.60	0.03	67.7	3 4 6	1.48	181	-
918	139641	R64231	0.12	0.31	2.67	1.98	7.39	1.13	6.10	2 6	92.04	-
ā	609228	AA179161	0.0	0.25	2.67	0.61	2.46	0.83	87.	2 .	<u> </u>	• •
8	3777	A00000A	0.58	10.	2.64	1.47	1.50	8.81	1.78	1.27	<u> </u>	
į	2000	Ne 2048	2	0.87	2.63	2.06	6.78	1.51	2.87	1.46	<u>R</u>	- ,
7	2007	01000	3	, c	2.83	2.25	0.11	2.32	0.83	1.71	8. SS	-
7	ercone !	A14/384		3 6		2.59	1,25	1.32	6.01	1.69	2.86	<b>,</b>
Z.	23023	R15580	9 .	ξ, <b>.</b>	70.7	2.25	2.58	2.24	8.64	0.96	<u>5</u>	-
824	273546	N33274	2.5	6.55	0.7	64.6	111	8.50	127	0.72	1.45	-
83	843049	AA485963	0.10	c7:0	10.7	1.35	2 19	2.32	6.51	1.78	1.46	-
858	594517	AA169645	32	81.32	2.60	0.5	2 6	4	2.78	1.77	2.03	-
128	453175	AA700218	0.99	2.58	2.60	5.15	2.24	50.	97.	0.37	0.70	-
828	360357	AA013355	0.26	0.67	2.80	1.94	3.92	07.1	9	72.4	1.53	-
828	345525	W72437	3.95	10.21	2.58	2.40	2.96	1.23	5.63 6.0	r 60		•
8	344274	W73811	0.13	0.33	2.58	0.08	0.69	12.71	0.53	0.28	2 6	- •
Ę	1001	AA733105	0.03	0.08	2.58	9.39	0.82	1.72	2.27	0.98	P. 0	- ,
8	787745	44418584	1.03	2.65	2.58	1.48	1.68	2.03	2.47	9.56	2.28	- ,
5	744777	AAR20355	0.86	2.22	2.58	0.98	2.48	5.63	1.61	1.81	88. I	
3 8	#40A7K	4444766	0.07	0.18	2.57	2.	1.73	1.89	1.42	7.19	1.57	<b>-</b> .
\$ 8	C70010	44872024	200	0.51	2.57	1.98	1.32	1.27	7.42	2.20	124	-
2 8	000/4	44,00063	2 6	12.6	2 67	1.20	1.20	1.67	8.80	0.59	1.88 88	-
8 1	201410	70704		1 5	2.57	1.13	1.1	1.07	9.0	0.72	10.42	-
	707	761047		3 ?		2.29	1.57	707	1.01	75.0	7.34	-
3	079054	WICIO/W	0.0	47.	, t	1.86	5.20	1.35	2.96	1.57	2.33	-
3	50/05	K72003	) ;		75.	1.75	2.23	0.74	2.28	2.59	5.66	-
3	20428	K42061	2 6	6 6	5 2	38	0.38	2.96	9.80	0.36	1.25	-
<b>3</b> 3	701.007	2/8/2	3 5	77.	2.53	2.42	5.39	1.41	2.09	1.48	2.38	-
¥ :	*10/07	75007N	8 6	2 2	3.83	1.78	1.91	1.19	1.88	8.28	0.17	-
3	2000	BL8//X	2 6	<b>5</b>	2.52	2.29	7.29	1.91	0.18	1.86	1.60	-
<b>:</b>	02/461	D2265	6.6	2 6	3.53	1 20	1.50	1.33	0.70	9.11	1.24	-
3 3	CORCE	AA/3/304		1 2	2.51	1.63	5.07	2.31	1.74	1.69	2.63	-
3 3	A 10000	2000		878	2.51	1.12	1.30	2.13	18.	1.86	0.82	-
Ì	74080	080081	2 5	0.45	2.50	1,45	2.91	1.52	1.12	5.41	2.57	-
3	592592	AA159577	3.36	8.37	2.49	1.59	0.54	212	9.45	0.77	0.49	-
•						``	28					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

		Gen Bank			Fold/AveTumor//	Fold(MPM120/A Fold(MPM120/A Fold(MPM121/A Fold(MPM124/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A # of Tumor	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A	Fold(MPM126/A	* of Tumor
	IMAGE	_	AVEBPH	AveTumor	•BPH)	VeBPH)	veBPH)	VeB(H)	veBPH)		:	
		Number			. :	1	1 24	2.42	1.57	1.07	1.62	-
980	591055	AA161097	0.08	0.20	2.49	00.7	19	38	1.52	1.97	131	-
951	267252	N24579	0.83	2.07	2.48	6.02	7.0A	5	108	1.21	9.12	-
252	53383	R16245	1:1	2.75	2.48	1.34	1.22	- 6	53	1.88	0.48	-
8	327058	AA284287	o. 16	0.39	2.47	9.38	1,43	2 6	2.81	3	1.48	-
2	284544	N69425	0.61	1.25	2.47	2.70	6.07	3 9	, v	98.1	1.92	-
ě	20000	HEADO	28.65	70.79	2.47	2.13	1.82	04.1		9	2,54	-
3 8	400,000	100074	72.0	0.67	2.47	1.83	2.00	1.57	<u> </u>	9 6	35	-
8 1	00.4311	78081	3,5	180	2.47	1.81	<b>1</b> .	1.23	1,48	9 7.0	8	-
924	8//019	4C/184V	;		2.48	1.61	5.16	2.27	1.48	177	9. 6	
928	431933	AA678164	= :	3 3	÷ ;	0.87	2.24	2.13	1.48	1.27	0.81	- •
88	359653	AA010872	0.32	0.80	2.40	(P)	2.61	1.59	5.29	1.80	5.03 5.03	<b>,-</b> ·
8	681865	AA258157	0.0	1.51	2.46	? .	37.0	10.09	1.06	0.89	1.07	-
<b>8</b>	239199	H70942	0.91	2.25	2.48	0.80	, q	0.58	0.87	1.05	1.89	-
862	66564	T67058	0.31	0.76	2.48	2.60	90.	2 6	1.18	2.82	5.77	-
883	215000	H73241	3.81	<b>3</b> .8 <b>x</b>	2.45	2.77	1.20	7 6		1.26	1.22	-
8	810773	AA481751	0.87	2.12	2.44	1,12	1.19	79.6		1 87	1.43	-
š	282000	N51107	0.53	1.29	2.44	2.00	1.90	5.73	2 5	1 37	98.0	-
3 8	200.00	2000	2.54	6.18	2.42	1.35	2.68	2.98	9.20	5:0	990	-
8 8	20102	7798677	710	0.40	2.42	1.55	1.08	<del>.</del> 2.	0.80	) T. F.	2 18	. 🖵
2		+1 C07+W		72.03	CF C	2.58	6.50	0.47	1.05	1.75	5 6	- •
88	887658	<b>W4</b> 86786	07.6	17.7		88	1.77	2.00	1.45	2.38	20.0	_ ,
8	111884	T84886	0.03	0.0	14.7	<u> </u>	149	2.43	5.19	2.25	1.48	-
970	744897	AA525784	1.69	<b>*</b>	2.40	(c.)	7.0	2.67	5.14	1.19	1.88	-
£	46411	H09164	3.30	7.91	2.40	1,72	97:1	. «	2.12	2.35	1.52	-
22	798781	AA460729	0.59	1.42	2.40	1.08	וכיר י	9 9	1 8	0.90	1.70	-
973	1031273		0.38	0.91	2.39	1.15	<b>5</b> .	40°L		08.9	2.65	-
478	46129		0.0	0.16	2.39	0.81	1.2	08.1	4. 6.	7.40	2.27	-
975	220389	H88816	1.1	2.65	2.39	1.00	1.09	9:1	. T	75	0.88	-
978	1292170	_	1.48	3.50	2.35	1.36	9.50	30.00	5 6	8.51	1.68	-
3	35.0711		0.78	1.88	2.38	1.32	2.39	7.0Z	0; i	7 78	28	-
	4308A	A A R 7 R 2 U 3	0.65	8	2.38	1.54	2.21	2.38	<b>3</b> 0.0	07.1	. K	•
0/8		201000	9.0	4.43	2.38	1.71	0.91	2.32	2.25	9/:	3	٠.
818	32575	108C*X	9 5	? .	237	0.80	1.53	1.02	8.43	1.41	7. C	- ,
960	<b>84</b> 079		20.0	3 5		8	0.90	0.69	0.41	10.29	1.05	-
981	1388395	•	0.30	7/0	16.3	96.5	102	8.38	2.45	0.70	0.28	
982	197793	R93729	0.78		2.37	86.1 9.64	1.43	1.12	6.66	2.63	1.71	<b>-</b>
S	26738	R39891	7.85	18.11	2.37	<b>\$</b> 0.0	•	!				
							29					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

		Gen Bank			Fold/AveTumor/Av	Foldiax. Tumodav Foldimpint20/A Foldimpint21/A Foldimpint24/A Foldimpint22/A Foldimpint23/A Foldimpint26/A # of Tumor	old(MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A F	old(MPM126/A	# of Tumor
	Clone ID	Accession	AveBPH	AveTumor	•BPH)	(NeBPH)	VeBPH)	veBPH)	veBPH)	veBPH)	veBPH)	Samples
ě			9	7	2.38	1.26	96.0	127	8.89	1.09	0.62	-
\$ 5	SADAS	43287	9 5	- 6	2.36	75.	0.57	2.63	1.43	1.70	0.30	-
200	CL89/7	N346Z4		7 7	2.35	6.87	2.92	0.89	0.84	0.87	1.75	-
0 P	20005	AA034113	2 4	. 5	2.35	7.31	1.59	1.65	1.23	1.13	1.21	-
	610171	280781	<b>1</b>	1.28	2.35	1.29	1.46	1.13	1.43	7.48	1.30	-
9 0	34442	011000	1 70	8	2.35	0.88	0.88	2.43	8.64	0.85	0.41	-
	747547	0,000,44	200	0.48	2.34	1.71	2.12	1.52	1.14	‡	2.13	-
26 6	74576	01000	2 6	0.70	234	1.23	1,45	1.56	5.33	1.78	2.68	
3 8	34/0/20	W423700	200	0.13	234	0.52	0.74	76.0	1.18	1.22	9.40	-
8	285114	120820	5 6	0.80	7.3	7.86	0.89	1.68	1.01	0.44	2.14	<del>-</del> ·
8 8		760060	9	3 0 %	7.5	1.72	1.97	2.52	0.59	0.96	6.28	-
į	260040	+2+LALVY	<u> </u>	3 5	233	5.38	0.72	2.23	4.	2.12	2.12	-
6	201403	79//9N	\$ 5	2 5	233	1.83	2.67	28.1	5.6	1.01	1.19	-
2	240318	CR/ROL	?	3 :		72.0	0.58	9.21	0.83	66.0	1.48	-
186	252684	Heecs	<u> </u>	2.02	 	80 -	17.	2.18	1.05	6.70	52	-
88 6	302180	W18724	<u> </u>		2.33	1.55	1.60	7.25	1.31	1.20	96:0	<b>-</b>
	130742	A A A 2004 4	2. 5.	305	2.33	0.67	<b>3</b> 9:0	0.61	9.75	1.41	7870	-
3 5	23467	4400044	80.0	0.85	2.32	1.41	1.54	1.91	1.41	<u>2</u> .	6.13	-
3 5	281241	HOR244	0.85	1.87	2.32	2.51	5.63	1.48	1.28	1.99	1.05	-
3 5	21030	A A 7 K B 4 7 D	9	0.13	2.32	5.12	1.82	2.46	2.71	0.86	0.85	-
<u> </u>	381008	AA001438	0.48	1.07	2.32	1.41	0.59	0.79	9.48	0.33	<del>7</del> .	-
į	807078	AA400785	2.16	4 89	2.32	0.83	1.17	8.26	1.19	1.30	1.05	-
3 5	3344	073634		90 0	2.30	<b>1</b> 8.	0.79	1.98	2.04	2.01	5.13	-
3 5	ALL NO.	44006228	0.25	98	2.29	0.96	0.54	0.17	2.06	9.31	0.70	-
3 5	1030738	AAAAAAA	7	1.23	2.28	1.63	1.43	7.27	1.2	1.36	0.83	-
Ş	005122	44497007	18.96	43.36	2.29	0.95	1.38	1.79	<b>9</b> .04	2.82	0.85	-
1010	305809	None	0.92	2.10	2.28	0.83	1.72	1.21	5.27	2.14	2.50	-
101	30502	R41685	0.11	0.28	2.28	5.01	2.84	2.00	1.62	122	1.18	-
1012	187794	44498570	0.68	1.56	2.27	0.80	0.89	1.89	1.85	6.37	1.85	-
10.	730082	AA034179	0.40	0.92	2.27	0.70	1.27	1.75	2.28	5.09	2.67	-
101	48238	H10865	0.17	0.38	2.27	5.57	1.4	2.71	1.32	0.42	2.18	-
1015	703739	AA278749	8.47	19.24	2.27	0.85	0.71	0.71	1.69	0.62	9.05	-
1018	342720	W88398	0.42	98.0	2.27	0.42	0.02	0.87	1.58	5.54	1.19	-
1017	25158	R38835	1.15	2.61	2.27	1.08	5.81	2.33	2.50	0.73	1.15	-
						<b>6</b> 3	30					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

ACCURATION				(HGRPH)	veBPH)	VeBPH)	veBPH)	Fold(AveTumorfAv Fold(Mrm) Lunch VeBPH) veBPH) veBPH) veBPH) veBPH) veBPH) veBPH)		
,			(HAR			3	<u>.</u>	0.47	6.64	÷
	6	ς.	2.26	1.64	2.38	L8:0	2	0.8	2.58	-
AA411607	90.0	2 5	300	2.09	0.54	1.89	5.13		0.17	-
AA055979	90.0	<u>.</u>		0.33	0.35	0.39	11.15	7 ; ;	4.	-
N31605	2.29	5.18	C7.7		1.65	0.93	6.21	2.11	2 3	
AA7777870	5.06	11.36	225	6.93	2	2.83	6.99	1.42	62.0	- •
H19068	0.37	0.84	2.24	1.38		4	1.26	8.84	0. 1	- •
AA778089	121	2.70	2.23	0.48	?	10	2.62	1.52	1.65	_
A A 2 842 83	0.0	0.21	2.23	0.18	67.1	103	1.27	1.26	1.20	
A & 0.19318	0.22	0.48	2.22	79.7	0.92	1.02	1.35	1.81	1.21	-
H10208	0.78	1.73	2.22	6.51	1.41	90:	40.50	1.07	2.87	-
87701.0	98.0	0.85	2.21	0.63	1.56	 4 A	1.73	1.36	1.27	-
AA478257	79,0	1.18	2.21	.60 .60	2.11	9 6	1.05	10.80	0.55	•
TK1220	600	0.20	221	0.08	0.07	0.70	1.80	2.	2.16	-
HORZER	000	0.07	2.20	1,59	5.46		9.15	0.27	0.27	•
A A 465495	10.76	23.71	2.20	99:0	0.35	76.7	0.71	7.63	1.13	•-
A A 77/67.55	0.48	1.05	2.20	1.24	1.48	70.	90	0.08	1.59	-
AA284278	0.25	45.0	2.20	1.76	\$. i	ē 8	5.87	1.47	1.30	•
T52320	4.01	8.81	2.20	0.86	1.78	76:	76.0	1.53	5.21	-
AA883170	0.57	1.28	2.19	1.91	2.76	<u>.</u> £	60.0	90.0	9.58	-
N54551	0.25	0.55	2.19	0.59	51.1	6 52	0.83	0.76	0.62	•
AA185512	1.86	4.07	2.19	0.77	0.03	68	1.69	1.34	5.45	•-
N73584	1.50	3.29	2.19	1.03	9. 5	. G	0.50	0.22	10.37	-
AA706835	0.16	0.35	2.18	0.69	0.71	35	<del>1</del>	1.27	1.87	-
AA459692	0.78	1.70	2.18	1.13	07.1	2 15	6.01	1.14	1.78	-
R88246	0.05	0.11	2.18	1.50	0.53	2 7	0.12	0.12	2.07	-
H09618	0.08	0.18	2.18	1.76	0.47	42.4	1.70	1.05	6.35	
W72838	0.42	0.91	2.18	1.28	) <del>(</del> ) (	980	1.07	0.55	57.0	-
H22858	0.42	0.91	2.18	8.97	0.79	9 7 8	0.19	0.86	0.19	-
N50808	0.05	0.11	2.17	10.72	0.90	1.25	0.62	6.45	1.19	-
AA890147	0.51	1.09	2.17	1.57	7.72	36.0	0.25	0.25	1.7	-
AA010223	0.04	0.09	2.17	2.12	2.37	1.37	1.17	5.78	2.04	-
H88312	0.70	1.52	2.18	1.47	7.17	92.0	0.85	5.42	2.24	-
R81831	1.73	3.73	2.16	2.42	07:1	1 55	0.49	0.57	9.51	<b>-</b>
W86075	0.23	0.48	2.16	0.42	0.03	1.24	0.68	0.70	<del>1.</del>	-
R62289	0.64	1.37	2.15	1.94	5.	<u>!</u>				
ì										

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	NAGE	Gen Bank Accession	AVERPH	AveTumor	Fold(AveTumorlav Fold(MPM120/A Fold(MPM121/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A # of Tumor 	Fold(MPM120/A I	Fold(MPM121/A	Fold(MPN124/A	Fold(MPM122/A veBPH)	Fold(MPM123/A veBPH)	Fold(MPM125/A veBPH)	# of Tumor Samples
		Number						30	1 70	6.30	0.82	- -
1052	131626	R23735	0.10	0.22	2.15	0.63	1.32	ca.		90	8 0	
1053	32683	R43319	050	1.07	2.15	0.99	1.42	1.28	05.7	9	) ·	. ,
Ş	1031717	A A B 100 5 8 4	1.03	2.22	2.15	1.83	1.78	1.80	5.23	1.14	D	- •
105	354708	N2777	139	2.87	2.15	99'0	0.43	9.65	0.73	0.79	0.60	<b>-</b> .
	700007	A 4 6 3 9 0 7	89	9	2.14	1.48	2.24	2.10	5.53	0.52	1.00	-
8 5	100001	7807540		100	2.14	0.31	5.65	0.31	2.51	1.42	2.64	-
	2000	107904		9	4,	10.49	8	1.10	0.61	0.20	0.38	-
	CARRE	0//4	9 5	2	<u>, , , , , , , , , , , , , , , , , , , </u>	131	0.35	<del>1</del> .	0.49	8.11	1.08	<b></b>
SCOL S	12840	AA282023	2 5	9 6	2 5	36.1	1.22	6.15	1.77	1.37	0.92	-
<u> </u>	208123	A010136	÷ ;	3 8	5.15	80.	148	2.05	1.23	1.34	5.57	-
<b>8</b>	20/00	H60361	7 6	<b>8</b> 6	2 5	5 6	5.03	1.05	1.68	1.36	1.40	-
28	10071	AA757764	9 :	7 :	7	3 5	53	1.70	1.68	1.15	6.41	-
1063	78946	T61782	3.51	<b>3</b>	21.2	C <b>7</b> -	56.0	00.1	1.25	1.33	5.19	-
<del>2</del>	42373	R67147	7.51	3°.5	11.7			68.0	1.69	0.70	6.69	-
1065	199628	R96598	0.18	0.37	2.10	) . (0.	) - - -	0.48	2.13	7.	6.32	-
100 8	29828	R42331	1,30	2.72	2.30	2.03	2, 5,		1.28	0.00	1.30	-
1087	809828	AA455521	0.68	.39	2.10	9.76	<u> </u>	8 0	1.09	0.95	7.10	-
1068	280561	N62376	2.80	5.88	2.10	.s. ;	Ö. 6	9 4	1.58	1.28	0.92	-
1069	781088	AA430002	0.62	1.31	2.10	1.23	96.0	0.0	5. t	1 17	8.21	-
1070	282384	N51987	0.51	1.08	2.09	1.05	66.0	08.	57.1			• •
1071	782873	AA447585	0.21	<b>7</b> :0	2.08	0.47	1.67	0.47	1.20	88.7	æ 6	- <b>-</b>
1072	73785	T54672	1.35	2.70	2.07	1.21	1.21	1,49	77.5	1.88	<u> </u>	- ,
1073	730124	AA412470	0.22	<b>\$</b> .0	2.05	2.93	1.29	0.05	2.00	5.24	0.73	<b>,</b>
1074	61403	H18924	0.28	0.67	2.05	1.36	0.99	0.32	1.09	7.73	0.79	-
1078	38497	R40124	0.34	0.70	2.05	0.64	1.05	0.28	0.36	9.67	0.30	-
1078	1048723	AA670647	0.71	<del>-</del>	2.03	1.67	1,45	1.18	1.05	6.22	0.63	-
ţ	141815	870685	8.27	18.77	2.03	0.45	0.67	0.71	0.56	8.86	0.91	-
1078	767.37	A 4.37009	128	2.60	2.03	09.0	96.0	0.58	2.60	1.86	5.64	-
0.70	244640	N27841	90	60	2.02	7.29	1.20	0.85	0.79	1.68	0.22	•
1080	428189	AAOO1878	0.20	0.41	2.02	0.05	0.31	0.75	0.65	89.68	0.68	<b>~</b>
Š	050470	A 4 500102	2.48	200	2.02	1.00	1.18	6.48	1.57	90.0	0.92	<b>-</b>
1 60	824044	A4491208	0	0.18	2.01	0.11	1.13	0.92	1.59	6.82	1.38	-
1083	1458721	AA864875	12.87	26.09	2.01	1.29	1.62	1.39	5.81	1.48	0.49	-
100	757156	AA443938	0.49	88.0	2.01	0.95	1.21	0.63	0.37	8.19	0.71	<b>-</b> -
1085	252830	H88281	1.63	3.26	2.00	1.12	0.93	7.24	0.89	0.87	0.98	<b>-</b>
							32					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

5.45 1.50 2.33 5.66 1.13 1.00 1.62 5.66 1.18 1.05 6.39 1.19	1.60 0.77 1.57 5.64 1.01 1.86 6.15 1.30 0.48 0.69 1.76 0.57 0.87 1.28 1.57 0.30 1.70 1.67 5.10 0.62
5.45 1.50 2.33	1.05 6.39 1.19 1.57 1.57 1.57 1.57 1.57 1.57 1.57 1.57
5.45 1.50 4.33	1.05 6.39 1.60 0.77 1.01 1.86 0.48 0.69 0.87 1.28 1.70 1.67
5.45 1.50 1.13 1.00	1.05 1.01 1.01 0.48 0.87
1.1. 1.13 1.18	
	1.06 2.00 1.33
2.00 2.00 2.00	
	1.98
4.18 50.73 1.48 4.66	15.78
*	6.03 72.0
	4 4 770020
24916 1837296 123539 1461664 1467207 78022 37665 462545	

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

		Gen Renk			F. 1885 T.	Fold(MPM120/A	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A	Fold(MPM126/A	s of Tumor
	IMAGE	Accession	AveBPH	AveTumor	Fold(Ave Lumor)Av •BPH)	veBPH)	VeBPH)	veBPH)	veBPH)	veBPH)	V•6PH)	and units
		Number			•	;	ć	102	5.71	<b>7</b> .	680	-
1120	243321	H94720	1.08	2.00	1.86	0.82	98.0	70.1		105	5.57	-
123	243803	N49725	9	1.19	1.86	1.16	1.11	<u>.</u>	2 7	1.87	0.17	,
		44.67457	90	0.11	1.86	1.70	5.30	2.13	0.17	5 6	2	•
7	2016	200		234	1.85	0.50	6.17	1,41	0.72	7.7	80.	
1123	80371	165765	17.1	, , , , , , , , , , , , , , , , , , ,		168	2.25	69.0	5.70	0.12	0.67	-
1124	731029	AA421269	0.0	er.o	6.	770	0.82	0.58	5.74	2.61	0.92	-
1125	284882	N66737	0.33	0.61	68.	<b>T</b> 6	7.0	1.52	7.50	0.73	0.26	<b>*-</b>
1126	1031009	AA635186	6.84	10.78	1.85	9.0	o e .		E	1.16	1.13	-
1127	1030637	AA608770	0.76	14.	1.85	1.20	1.32	3	5 + 5	9.9	0.07	-
1128	273168	1979KN	0.15	0.27	<b>2</b> 8.	1.65	0.62	<b>8</b> 5	1 6	080	1.28	-
1120	66317	TB6616	8.85	16.25	<u>.</u> 2	6.02	2.40	0.48	7 7 7	08.0	0.92	-
		70000	=	203	1.83	98.0	0.98	0.40	3	9 6	90	•
1130	28507	40077V	= :	3	4.82	0.63	0.91	5.53	<u>.</u> ‡	1.09	90.	- ,
1131	487088	AA045300	2.58	PO I	7	9	0.87	5.81	0.92	0.75	1.68	-
1132	253598	H89376	1.38	5.60	18.1	0.00	1 47	5.27	1.26	1.20	0.61	-
1133	1056186	AA621001	0.86	<u>4.</u>	1.79	0.96	÷ :		843	0.98	0.62	-
133	1806837	AA996026	31.85	67.14	1.79	1.13	200	3 .	9 6	0.17	1.77	-
1135	243731	W89213	90.0	0.11	1.78	0.21	5.41	00.7	9 6	080	0.12	-
	25058	708870	0.09	0.15	1.78	0.12	0.55	1.10	78.7	<b>8</b> 9		•
2 5	2007	70000	2,50	4.78	1.78	0.89	1.40	1.21	5.01	1.13	3.	- +
35	100/07	<b>5</b> 0000		9 6	1.7	0.60	9.0	6.31	0.75	1.01	51.13	- •
1138	128006	K09729	2.6	8 6	2.38	0.81	0.87	1970	0.83	6.53	0.83	-
1139	397804	AA701034	0.18	0.32	2 :		9	0.88	0.50	0.50	6.13	-
1140	42271	R60328	0.05	9.0	1.70	90.7	3.5	1.28	0.30	0.30	1.28	-
1141	71674	T67920	0.24	0.42	1.78	0.0	\$ 7	7.0	2 19	ş	90.0	-
1142	34084	W57616	0.17	0:30	1.74	0.40	F8:0	;	9	2.82	0.10	-
1143	788258	AA454098	0.10	0.17	t.1	0.18	2.20		2 6	, C	5.93	-
1717	142397	R69934	90.0	1.13	1.72	06:0	.8	8.	78.0	9 6	1 28	<del>-</del>
477	ARAN2	140477	0.89	1.53	1.72	0.48	0.39	0.57	<b>4</b> .9	07:1	2 7	
		988	0.47	0.80	27	0.96	0.75	1.23	5.01	3.56	0.7	- •
9 !	3007	20000		0.00	22	0.86	60.0	0.31	0.90	6.93	1.23	
414	0/404	5/00/50/5	: ;		2.	117	2.30	5.68	0.97	0.00	0.00	-
1148	2440	AA453610	9 6	3		7	5.02	1.21	2.24	0.85	0.51	-
140	471408	A035317	90.0	<b>4</b> .0	<u> </u>	6.4	5	1.30	5.72	0.93	0.3 £	-
1150	745001	AA626011	<del>.</del> .	2.45	<u>.</u>	0.67	8:-		0.88	6.09	0.78	-
1151	609155	AA176867	2.10	3.59	1.7	0.49	<b>\$</b> ;	80.9	2.50	0.72	0.72	-
1152	796549	AA480274	0.01	0.05	1.70	0.72	<b>3</b>	90.6	000	1 8 Y	1.23	-
1153	363003	AA018866	0.40	0.68	1.70	1.05	1.03	0.60	0.60	9	<u>!</u>	
							8					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

					Softwork Entringm121/A FoldINPM124/A Fold[MPM122/A Fold(MPM123/A Fold(MPM123/A Fold(MPM125/A FoldINPM125/A Fold	<b>V</b>	End/MPM121/A	Fold(NPM124/A	Fold(MPM122/A F	old(MPM123/A	Fold (MPM126/A	# of Tumor
	INAGE	Gen Bank Accession	AVBBPH	AveTumor	Fold(AveTumor/Av	FOID(MFM12WA VOBPH)	veBPH)	veBPH)	veBPH)	VeBPH)	(u.co.)	
	Clone ID						0.44	1.28	0.83	6.01	0.14	<del>-</del>
1355	271483	N31508	20.0	0.12	1.69	1.75	<b>t</b> 9	1	5.86	0.92	0.76	<del>-</del>
7	48.158	H09838	2.53	4.28	1.69	0.88	0.50	78.0	1.35	0.70	6.17	-
2 2	8754R1	A A 504354	1.09	1.83	1.69	1.60	0.45	1 2	0.41	0.91	5.19	•
8 1	4204060	AA707460	250	0.55	1.89	1.71	1.08	5 6	0.82	0.78	0.50	-
130/	201871	Wenser	98.0	1.80	1.66	0.87	0.68	200	, v	9.0	0.62	<b>-</b>
2.5	433358	ROARSO	0.61	1.01	1.68	1.52	0.76	1.50	<u> </u>	5.30	0.16	-
	20000	NORTAG	900	0.10	1.65	0.27	2.45	6.39	23	99.0	5.01	-
2	4203440	_	0.46	0.76	1.65	1.60	1.37	20.1	1.09	7.12	0.60	<b>-</b> -
1167	78953	_	2.45	4.03	1.85	0.50	0.28	6.53	0.23	0.16	7.41	-
1107	772918	AA479912	90.0	0.10	1.64	0.16	1.15	0.73	47.0	5.35	0.35	-
3 2	24889	R43017	0.23	0.37	1.62	2.56	0.27	2.50	75	0.97	6.3 3	-
185	84141	171042	9.0	0.02	1.62	2.59	8.76	£ 53	5.01	2.20	0.97	-
¥ 5	969638	AA663592	1.87	3.03	1.62	0.64	0.38	C +	7.87	0.18	0.34	-
1487	810697	AA457686	0.14	0.22	1.61	0.07	0.07	 	5.00	0.14	89.0	-
	37234	R35283	0.07	0.11	1.61	1.33	1.88	0.00	5.63	0.11	2.13	-
1180	186132	H39580	0.24	0.38	1.59	<del>.</del> 9.	0.41	72,	0.62	5.54	0.10	-
170	112658	T91168	0.10	.0.16	1.59	1.13	C. 0.	07.0	0.96	6.69	0.09	-
171	788117	AA460961	0.11	0.18	1.58	0.29	1.78	900	0.11	0.11	5.73	-
1172	41321	R56906	90.0	0.14	1.57	1.21	1.33	030	1.31	0.75	6.14	-
Ť.	429448	AA007819	0.30	0.48	1.57	0.28	0.63	181	60.0	60:0	99.0	-
1174	882571	AA878515	0.11	0.17	1.65	0.89	08.0	27.0	0.61	0.40	5,41	-
1175	448813	AA777590	<u>1</u>	1.80	1.55	1.72	0.40	97.0	6.22	0,22	0.33	-
1178	40358	H15104	0.22	0.34	1.54	0.35	1.37	0.10	0.86	6.51	0.31	-
1117	810891	AA459519	920	0.55	1.52	0.56	9.50	5.42	0.73	0.70	0.66	-
1178	694123	AA189475	<u>2</u> .	2.89	£.	0.83	27.0	0.59	0.31	0.31	0.31	<b></b> -
1179	869375	AA679907	9.03	0.05	1.51	. C. C	64.0	0.11	0.88	0.62	0.27	-
1180	128248	R11510	000	0.14	1.50	S 6	200	77.0	0.51	5.78	0.55	-
1181	22895	R38640	0.09	0.14	1.50	10,1	9.0	5.33	0.81	0.65	0.71	
1182	286404	N87274	<b>5</b> .	222	1.48	0.77	0.02 0.03	0.73	1.3	5.61	0.31	-
1183	366915	AA027147	0.16	0.23	<b>4</b> .	0.60	9 6	1.35	0.20	0.25	0.25	-
18	591465	AA160685	<b>5</b> 00	0.08	1.42	0.40	5 6	0.24	0.82	44.0	0.63	-
1185	344595	W73834	0.37	0.52	1.41		0.92	0.31	0.31	0.84	6.29	
1188	781110	AA430042	0.03	<b>5</b>	7.38 5.50	0.45 8 t C	0.20	0.98	0.73	0.24	5.97	-
1187	291690	N87832	0.0	<b>0</b> .08	<b>8</b> .∓	5						
							35					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

Fold/AveTumoriav Fold(MPM126/A Fold(MPM124/A Fold(MPM122/A Fold(MPM123/A Fold(MPM125/A B Of 1umorian Fold(MPM126/A	0.49	0.07	5.71	0.46	0.91	0.50	0.82	0.45	5.44	0.33	5.26	5.27	
Fold(MPM123/A F	3,5	0.07	0.11	0.88	0.27	0.18	5.08	0.29	74.0	0.12	0.36	0.12	
Fold(MPM122/A	0.42	1 85°C	0.75	0.72	91.0	0.78	0.43	5 7 3	0.43	0.12	0.61	0.67	Š
Fold(MPM124/A	5		0.20	00		· •	9 4	8.0	0.20	97.9	4,0		0.00
Fold(MPM121/A	;	0.52	0.78	0.35	0.46	18.6	0.29	<b>3</b> .	0.45	0:00	0.12	A (	0.12
Fold(MPN120/A	•	0.74	6.50	09.0	0.52	0.60	5.83	0.43	0.45	0.29	0.12	<b>X</b> .0	0.12
Fold(AveTumor/Av 1		1.37	1.37	1.35	1.33	8.	1.31	1.29	1.27	1.25	1.21	1.19	1.18
AveTumor		<del>1</del> .	0.20	0.11	2.37	0.08	0.07	1.83	3.64	0.22	0.10	0.19	0.10
AWBPH		1.05	0.14	90'0	1.78	70.0	90.0	1.41	2.86	0.17	8	0.16	0.09
Gen Bank Accession	Number	AA670390	AA875953	AA129677	AA046430	R60722	H28216	AA688959	N95226	AA055491	AA160670	W39343	AA131516
MAGE	Close (C)	878469	1482258	665403	488207	42118	52594	854401	307337	377296	592728	322617	503545
		1188	1189	25	1191	1182	1183	2	<del>2</del>	28	1107	1108	1189

Table 2-2: T3NO vs. BPH: Gene upregulated 5 fold or higher in at least 2 tumor samples or 3 fold or higher in at least 3 tumor Samples.

Table 2-2: T3NO vs BPH: Genes upregulated 6 fold or higher in at least 2 tumor samples or 3 fold or higher in at least 3 tumor samples.

											A/CC+140401051-2	Powiuput23/A	Fold(uPut26/A	# of Tumors
	MAGE	Gen Bank	Tiagre	Secreted/Me	AvaBPH	AveTumor	Fold/AveTumoria Fold(MPM12NA Fo	Fold(MPM120/A	Fold(uputzila vsBPH)	FOIG(MPM124A veBPH)	VeBPh)	veBPH)	VeBPH)	Samples
Š	Clone ID	Number	Distribution	Bound						7.36	52.32	22.54	10.36	•
-	786180	AA453310	prostate		<del>2</del> .	133.47	<b>3</b>	140.63	188.10	24.6	93.00	3,53	35.80	•
	646037	AA196979	prostate		2.38	14.00	<b>\$</b>	85.24	8 5	<b>1</b> 2	526	41.24	18.50	•
· •1	284701	N64840	prostate	>	142	73.10	8	31.27	5 5	3	16.57	45.42	20.45	•
•	450049	AA703396	prostete	<b>\$</b>	245	78.62	8	28.52		90.01	10.72	21.45	63.61	•
40	245768	N55286	•		5	62	ឌ		3 6	22.40	15.57	11.81	13.20	•
•	231003	NSOBBO	3		24.05	<b>4</b> 23.66	₽ !	2. 2. 2. 2. 2. 2. 2. 2. 2. 2. 2. 2. 2. 2	2 7	8.28	10.57	5.71	721	•
1	136605	R35051		<b>&gt;</b>	97	<b>4</b> 3	<u>1</u>	8 67.02	# # *	6.15	824	10.23	5.96	-
•	00,000	R28307	prostate	>	<b>3</b> .	25.46	<b>+</b>	2 3	5 5	3.85	15.7	3.32	7,01	•
•	899903	AAA67236	prostate		ž	48.10	<b>5</b> .	8 5	3 7	11.99	10.01	17.00	10.66	•
2	170753	H5154	- Sa		10.0	0.12	<b>2</b> 2 (	97.	15.01	11,83	6.14	127	15.03	•
=	1000656	AA621761	•	<b>&gt;</b> -	3	12.57	₽ (	R P	2	8.58	7.12	8.56	1924	•
2	52.63	H20747	•		707	19.56	<b>a</b>	9 5	9, 45	7.85	4.72	3.65	53	•
2	300000	N91003		<b>&gt;</b>	8	Į ;	<b>.</b>	5 0	9	100	153	12.71	6.41	•
2	196975	R91396	•		કુ	2		200	50.05	989	7.18	5.75	8 20	•
15	21931	172543	Draws		<b>8</b>	1.87	~ •		101	7.14	10.04	4.07	5.15	•
=	792217	N79180	<b>pove</b>	••	0.0	0.42			1	. S. 33	5.13	6.23	11.57	-
=	625450	AA504784	•	Y/A	<b>1.</b>	12.18	ь (		7	8.60	4.79	4.07	5.5	•
=	425816	AAD04881			0. 0	28.5	۰ •	, o	10.0	6.21	7.48	3.67	8.48	•
=	1668	R86764			8	4.0 4.0	ь -	121	19.6	3.89	96.8	3.63	308	•
2	359701	A011178			29	12.8/	• •	117	8	4.16	3.84	3.52	3.85	•
<b>7</b>	243641	N49548		Maybe	<del>.</del>	3 3	• •	2.1	£.13	3.55	4,67	3.77	<b>4</b> .01	•
#	262035	H90608	•		g	8 8	• \$	15.20	2.46	18.94	12.53	10.40	100	<b>.</b>
n	267628	M62096			9		3 <b>\$</b>	290	28.16	14.54	07.71	<u>±</u>	3.92	× ,
*	201.00	AAA32103	prostate		15.97	7.1	<u>.</u>	191	24.39	10.24	5.27	13.26	1.76	•
ដ	258374	N31962	prostate		<b>R</b>	20.5	2 9	37.6	90	3.	3.06	33.80	<b>4</b>	•
*	210620	H67712	•		<b>2</b>	7	2 •	10.67	97	7.67	13.10	121	327	<b>5</b>
22	193923	RE3407	Į.	>	17.30	130.73	• •	970	27.78	3.78	3.02	177	350	•
2	37980	R01372	2		3 !	* *	• •	2.69	10.96	6.72	12.04	2.83	207	•
R	1001599	AA609485			7	7 7	• •	7.38	5.10	8.4	15.83	7.96	67.4	•
ጸ	86180	172338	Pop.		5 5	2 :	. •	92.8	3.60	13.77	2.12	4.36	6.18	<b>o</b> '
Ä	2007	NZBBBB	prostede	;	à		. «	50.5	10.82	5,48	10.02	ā.	3.81	<b>.</b>
z	462468	AA699685		ş :	<b>8</b>	9 %	. •	101	3.57	4.27	14.38	70.0	ą.	<b>.</b>
z	806276	AA663160	•	<b>≨</b> :	9	\$ ;	> «	0.12	4.48	4.71	8.	2.69	8,45	•
7	811562	AAASAS97	DOM	<b>&gt;</b>		1 3		5	4.65	17.7	5.71	11.11	7.88	ń
2	428438	AADO4412	•		3	3.12	ъ,	2 2	9	6,30	<b>8</b> .38	8.0	2.00	10
8	130138	R20613			700			2 88	4.87	4.10	1.	3,72	11.48	w)
37	347586	W81432	7		5	3			8	590	11.83	6.34	5.73	<b>16</b> 0
8	785729	AA460291	unspectied		200	a ii	D	***	<u>}</u>					

Table 2-2: T3NO vs. BPH: Gene upregulated 5 fold or higher in at least 2 tumor samples or 3 fold or higher in at least 3 tumor Samples.

	IKAGE	Oes Bank	Tlascue	SecretadiMe	Haber	AveTurbor	Fold(AveTumoriA	FOR(MPM120/A	FoldAveTumoria Fold(MPM120A Fold(MPM121A Fold(MPM122A Fold(MPM121A Fol	FOLKIMPM124/A	Fold(MPM122/A	Fold(MPM123/A I	Fold(MPM12&A	f of Tumors Semples
į	Close 10	Number	Clatribution	Bound			(HARRA	TI DO				č	5	€7
;			į		5	15.0	•	3.24	861	2.85	6.33	5		
8	200	AA606117	overy		3 3		•	2.31	3.63	6.60	3.01	13.00	3.10	o :
\$	998	H28290	Ş		3 3	3 5		2	29.52	3.68	4.21	<b>8</b> .00	<u>.</u>	<b>s</b> n
Ŧ	417078	W87823	•			3	, u		10.29	3.67	2.30	7,50	4.13	**
ą	157075	AA486071	•	<b>&gt;</b>	ŗ.		n •	196	198	5.92	808	2.79	177	•
â	241821	H03381		××	8	3		2 %	58.0	194	7.80	4.62	2,43	•
1	28062	R40970	prostate		3		n •	3 5	90	3.89	4.46	87.8	5.11	•
3	124128	R0134	•	>	<u>2</u>	2	<b>5</b> 1	3 :		986	11,43	4.02	4.20	•
\$	234621	H65030	yarot		Ş	13 13	wn	3 5		3	7	3.53	2.63	•
13	1278	R28287	colon		870	3	•	20.0	87		1	3.82	5.12	•
\$	287745	N82344	Pass		Ē	18.41	vo	8 7 F	3 3	975	3.79	6.37	35,7	•
9	271141	N34482	,		100	8	40	0.70	2	-	2 10	7,0	1,47	•0
8	1240414	AA788780			0.79	3.60	wo	8	7	9 0	92	3.20	3.63	wo
2	307249	NECACION	lests		3.36	16.26	en	9.25	FC.8	3	76.7	17.	3.	€0
2	763648	AA443688	OVE.		0.13	97	•	4.82		6.43			235	•
2	0077720	AA598982	, g		27.0	£,3	•	3.68	3,50	4	2 5		3	
2	283070	N61297	-		870	0.12	•	3.87	8.0	3.48	9.0g	<u> </u>		, wa
: :	<b>365008</b>	N21081	í		900	070	•	g. i8	3.28	322	8	8 3		
8 5	7	7000	į		19	36.37	4	6.16	7.80	320	3.85	<b>7</b>	2.	
8 5	***			>	940	8.	•	2.15	7.00	3.45	3.00	6.14		
à :				- >	=	4.62	•	3.40	4.63	2.64	1,	5.85 5.85	3.0	<b>.</b>
3 :	2074	1970	2	-	2	81.0	•	3.25	3.83	4.50	3.60	4.60	12.	ρ,
2		79995X			3 5	2		4.70	6.56	3.16	3.79	3.49	Z	0
8	2000	AA011006	e e e e e e e e e e e e e e e e e e e		1 2			3.10	3.36	3,88	99.4	4.99	0.89	<b>≈</b> n
=	126230	R06307	5	;	3 5	2 5	, ,	50.5	3.20	3,51	3.6	3.10	<b>5</b> 4.	•
62	375062	AA032221	prostete	<b>&gt;</b> }	8 ;	2 2	, ,	18.80	99.1	69.34	105.24	0.80	19,00	•
3	1466260	AA884926		ž	20	8.7	5 8	20 78	17	20.56	98.0	23.62	1.28	<b>~</b>
2	743628	AA634378	•	<b>₹</b>	900	17.80	<b>3</b> !	2007	48.43	3.22	1.15	1.7	1.51	•
2	1390664	AA843502	prostete		<u> </u>	8	<b>2</b> ;		180	1 1	87.09	15.40	1.72	•
8	12	RSTTPA	<u> </u>		9	3	<u> </u>	2	5	88	6.52	49.21	8.28	•
<b>:</b>	74.02	AA825890	prostate		<u>.</u>	7	<b>2</b> :	2 5	2 %	228	9.20	2.33	1.9	•
3	1884.73 17.	AA778728	prostate	ž	5	2	= •		97.00	13.60	13.97	1.00	8.	-
2	40165	A045074	color		9	3			51.6	25	1.31	14.4	2.70	•
2	1466424	AA662992	Press		<b>3</b>	3	<b>.</b>	2.5	2 6	33	24.30	4.57	6.15	•
=	380184	A013260			9	5	•	5 5	2 5	6.6	10.46	<u>-</u>	2,63	•
22	141740	R76229	etaleorq		0.18	121		201	200	183	2.85	10.25	15.67	•
ß	24916	R38066	•		80	9.6	0	300	; ;	***	9	2.38	3.28	•
7	581099	AA185878		>	9.70	80.24	•	9.37	72.7	<b>9</b> .7	2 0	7.	10.42	•
92	1050172	AA820996	muscle		11.01	70.00	•	8	90,4	Ę,				•
2	782383	AA31407			58.0	5.56	•	885	11.77	5.12	5.02	18.1		
2 5	1001113	AA600014	•		00	0.0	•	16.58	0.83	4,0	17	28	2 1	•
: #	416	AA70675e		Y/V	70	1.86	'n	2.60	6.47	15.9	<b>5.</b>	6.07	2.77	•
. 1	78.30	4444756	}		920	3.00	•	3.83	6.17	7.31	8.02	179	797	•
2 ;				>			· vc	6.20	20.0	6.65	5.30	0.28	3. 3.	<b>+</b>
2		8/776WW	e B	•	2	š	•	}						

Table 2-2: T3NO vs. BPH: Gene upregulated 5 fold or higher in at least 2 tumor samples or 3 fold or higher in at least 3 tumor Samples.

Gen Bank         Tissue         Profession         Outsituding         Bound         AvaBPH           AAA70036         AAA270036         Done         0.002         0.002         0.002           AAA3077         bone         0.002         0.002         0.002         0.002           NA40038         nypothalamus         0.002         0.002         0.002         0.002           AAA30318         prosisie         0.004         0.002         0.002         0.002           AAA30318         prosisie         0.004         0.002         0.002         0.002           AAA30319         prosisie         0.004         0.002         0.002         0.002           AAA30319         prosisie         0.004         0.002         0.002         0.002           AAA30312         prosisie         0.004         0.002         0.002         0.002         0.002           AAA77002         brain         V         15.01         0.002         0.002         0.002         0.002         0.002         0.002         0.002         0.002         0.002         0.002         0.002         0.002         0.002         0.002         0.002         0.002         0.002         0.002         0.002         <	March   Color   Colo				a Milita			Approximation of the	Fold/MPM120/A	Fold(MPM121/A	Fold(MPM124/A	FOM(MPM122/A	Fold(MPM123/A F	old(MPM126/A	# of Tumors Samples
ALABORS         Done         0.02           AAABOUT         bone         0.04           AAABOUT         bone         0.02           H08463         muscle         0.02           H08463         muscle         0.02           AAABOTS         bone         0.02           AAABOTS         bones         0.02           AAABOTS         bresst         0.02           AAABOTS         bresst         0.02           AAABOTS         colon         1.70           AAABOTS         colon         1.245           AAATTROZ         muspecified         NA         2.47           AAATTROZ         bresst         0.02           AAATTROZ         bresst         0.02           H01784         bresst         1.41           AAATTROZ         bresst         0.04           H030025         bresst         1.44           AAATTROZ         bresst         0.05           H030026         bresst         1.44           AAATTROZ         bresst         0.05           AAATTROZ         bresst         0.05           AAATTROZ         bresst         0.05           AAATTROZ         <	Avassam         Distribution         Bound         0.02           Avassam         bone         0.04           15339         bone         0.02           15203         hypothiacmus         0.02           15233         bread         0.02           15233         bread         0.02           15233         bread         0.02           15233         bread         0.02           15234         bread         0.02           15237         bread         0.02           15237         bread         0.02           15237         bread         1.71           15237         bread         1.41           15234         bread         1.41           1534         bread         1.43           1534         bread         1.74           1534         bread         1.74           1534         bread         1.44           Avazozzzz         bread         Y         1.72		Gen Bank	•	mbrane	AveBPH	AveTumor	VeBPH)	VeBPH)	veBPH)	veBPH)	(Ladia)		,	
ALANDESS         CORD         CU10         S         127         138         482         128           ALASSOTT         Done         CORD         CU10         S         128         319         482         128           ALASSOTT         Done         CORD         CORD         CORD         S         128         319         482         128           ALASSOTT         Innect         CORD         CORD         CORD         CORD         CORD         128         319         344         482         128           ALASSOST         Innect         CORD         CORD         CORD         CORD         447         482         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483         483	March   Marc	0	Mimber		Bound				ļ	6.67	0,33	1,52	3.	9.17	• •
7.5330         Done         0.04         0.18         6         3.22         3.17         2.03         1.08           7.5330         Done         0.04         2.17         5         3.18         3.54         0.09         1.08         5         3.18         3.54         0.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09         1.09 <td>  1,000,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000  </td> <td>_</td> <td></td> <td></td> <td></td> <td>0.02</td> <td>0.10</td> <td>vs.</td> <td>977</td> <td>2 5</td> <td>4.92</td> <td>2.90</td> <td>1.73</td> <td>3.26</td> <td>-</td>	1,000,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000   1,000	_				0.02	0.10	vs.	977	2 5	4.92	2.90	1.73	3.26	-
VAMESCRIPT         CORD			A44/9030	. ;		500	0.16	\$9	3.02	2	9 0	8.52	30	80.	•
National	1,55,529   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,59,54   1,5	_	AA68CO77	mon .		. 200	90'0	10	1.68	200	<u> </u>	14 80	62.0	2.97	•
New			153389	Pool		2	11.34	vs.	3.22	E (F)	3 1	5	270	10.45	4
National	Page		N68465	musch		3	2 47	w?	3.19	Z.	8	3 :	15	11.65	4
Page	Augusta         Code		N74392	<u>\$</u>		\$ 1 5	. 9	wn	3,44	0,63	4.63	7		5.71	4
AMSZB11         Insert         COCK         COLK	Avolgostis         based         0.03         1.03         6.73         1.03         6.73         1.03         6.73         1.04         7.14         5.35         1.04         7.14         5.35         1.04         7.14         5.35         1.04         7.14         5.35         1.04         7.14         5.35         1.04         7.14         5.35         1.04         7.14         5.35         1.04         7.14         5.35         1.04         7.14         5.35         1.04         7.14         5.35         1.04         7.14         5.35         1.04         7.14         5.35         1.04         7.14         5.35         1.04         7.14         5.35         1.04         7.14         5.05         1.04         7.14         5.05         1.04         7.14         6.05         1.04         7.14         6.05         1.04         7.14         6.05         1.04         7.14         6.05         1.04         7.14         6.05         1.04         7.14         6.05         1.04         7.14         6.05         1.04         6.04         7.14         6.04         7.14         6.05         7.14         6.05         7.14         6.04         7.14         6.05         7.14         6.05			hypothelemus		200	500	v	1.99	4.80	2.74	878		123	•
AvASE225         Deset         O.27         1.28         5         5.53         10.10         4.41         8.5           AvASE218         preside         6.65         6.47         5         5.53         10.10         4.41         8.5           AvASE218         presides         0.06         4.43         5         5.52         5.50         0.58         3.87         0.58         3.87         0.58         3.87         0.58         3.87         0.58         3.87         0.58         3.87         0.58         3.87         0.58         3.87         0.58         3.87         0.58         3.87         0.59         0.58         3.87         0.58         3.87         0.58         3.87         0.58         3.87         0.58         3.87         0.58         3.87         0.58         3.87         0.58         3.87         0.58         3.87         0.58         3.87         0.58         3.87         0.58         3.87         0.58         3.87         0.58         3.87         0.58         3.87         0.58         3.87         0.58         3.87         0.58         3.87         0.58         3.87         0.58         3.87         0.58         3.87         0.58         3.87	Avaigned by Market State			Teer		8	0.24	, v	1 92	6.73	7,83	3.55	9 :	6	4
Avvestories         pages         45,47         5         54,69         43,93         6,48         4,48         6,48         4,48         6,48         4,48         6,48         4,48         6,48         4,48         6,48         4,48         6,48         4,48         6,48         4,48         6,48         4,48         6,48         4,48         6,48         4,48         6,48         4,48         6,48         3,48         3,48         4,48         6,48         3,48         3,48         3,48         4,48         6,48         3,48         3,48         3,48         4,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48         3,48	Avversity         problem         686         4547         5         544         485         638         751         130           Avversity         color         100         104         5         504         485         538         751         130           Avversity         color         170         124         5         506         1031         134         135         130           Avversity         color         170         124         124         5         506         533         131         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134         134 <td></td> <td></td> <td>breest</td> <td></td> <td>071</td> <td>1,28</td> <td>n '</td> <td></td> <td>10.10</td> <td><b>1</b>77</td> <td>8.26</td> <td>9.7</td> <td>ă i</td> <td>•</td>			breest		071	1,28	n '		10.10	<b>1</b> 77	8.26	9.7	ă i	•
Available         Cold	AVATORIZE         COLOR         610         5         OLD         155         344         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         155         1			alatamar		9.6	45.42	so.	C7:C	27	6.36	7.51	55	90'	, ,
AMATTREZ         Colon         170         794         5         350         150         150           AMATTREZ         Polaria         112.6         152.7         55.7         5.90         5.30         15.8         5.50         5.30         15.8         15.8         15.9         5.30         15.8         15.8         15.9         15.8         15.9         15.9         15.9         15.9         15.9         15.9         15.9         15.7         15.2         15.9         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7         15.7	National State   Nati	_				800	o.10	<b>5</b> 0	0.48	3 9	98.5	3.38	1.85	240	•
MATTOR         Probability         12.65         92.29         5.52         3.54         3.57           MATTOR         Maybe         0.94         4.15         5         4.04         3.24         3.07           MATTOR         Maybe         0.94         4.15         5         4.04         2.47         3.77           MODAGE         unspecified         0.47         2.07         4.15         6.43         5.18         2.47         3.77           MODAGE         unspecified         1.41         0.20         4.15         6.45         5.19         4.43         3.77           MODAGE         unspecified         1.41         0.20         4.15         6.45         5.19         4.43         3.77           MODAGE         unspecified         1.41         0.20         1.27         6.43         5.19         4.22         3.77           MATTOR         1.72         0.15         0.20         1.27         6.43         5.16         4.23         3.77         4.22         4.22         4.22         4.22         4.22         4.22         4.22         4.22         4.22         4.22         4.22         4.22         4.22         4.22         4.22         4.22	AMPLIANCE         Colored State         1246         92.9         5.52         3.54         3.54         6.29         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34         2.34 <td>_</td> <td></td> <td></td> <td></td> <td>2.</td> <td>7.0</td> <td>un</td> <td>3.60</td> <td>lg Ot</td> <td></td> <td>1.01</td> <td>1,63</td> <td><b>4</b>.8<b>6</b></td> <td>•</td>	_				2.	7.0	un	3.60	lg Ot		1.01	1,63	<b>4</b> .8 <b>6</b>	•
MATZBATZ         PROMISE         NA         247         1122         6,485         3,486         3,297           HB1736	AATTROZ         Provides         NA         247         1122         5         4.85         3.84         3.84         4.81         3.84         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81         4.81 <t< td=""><td></td><td></td><td>58</td><td></td><td>12.85</td><td>59.29</td><td>\$</td><td>5.52</td><td>or o</td><td>3 7</td><td>95.0</td><td>238</td><td>2.87</td><td>•</td></t<>			58		12.85	59.29	\$	5.52	or o	3 7	95.0	238	2.87	•
MAT7802         MAT7802 <t< td=""><td>AVATYRO21         NAM         CAST         1,04         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14</td><td>_</td><td></td><td>prostate</td><td></td><td>3 1</td><td>**</td><td>**</td><td>4.85</td><td>3.66</td><td>3.87</td><td></td><td>187</td><td>3.79</td><td>•</td></t<>	AVATYRO21         NAM         CAST         1,04         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14         1,14	_		prostate		3 1	**	**	4.85	3.66	3.87		187	3.79	•
Helly-36	Helitide	_			*	7	1	· •	30.7	2.42	10.26	<b>e</b> :		344	•
W02256         unspecified         0.64         4.15         5.34         6.46         6.59           R88915	W02250         unspecified         OS4         415         5.34         6.46         6.53         1.35         557           R88015         unspecified         O47         207         4.15         4.21         1.34         557           R88015         unspecified         141         6.20         4.15         4.21         1.35         3.56           AVXO0X52         unspecified         144         19.15         4.21         5.19         4.21         1.35         3.56           H2324         bail         8.21         4.27         3.19         4.27         1.31         1.31         3.56           H2324         bail         8.21         4.27         3.19         4.27         3.19         3.11         1.81         1.81         3.14         4.27         3.18         3.14         4.29         4.29         4.29         4.29         4.29         4.29         4.29         4.29         4.29         4.29         4.29         4.29         4.29         4.29         4.29         4.29         4.29         4.29         4.29         4.29         4.29         4.29         4.29         4.29         4.29         4.29         4.29         4.29         4.29         4.2				Maybe	8	CC.4	•	1.85	2.47	3.73	9. 9.	20.0		•
R88815         Lodge of the color         207         217         643         771           AVOO426         unpedied         1,41         620         4,37         3,16         4,22         1,21           AVOO426         brash         4,42         19.15         4         3,16         4,37         3,16           H12254         brash         Y         1501         82.10         4         3,16         4,37         3,16           H43260         brash         Y         1501         82.10         4         1,97         3,36         4,39         2,50           H43300         brash         Y         1501         82.10         4         1,97         3,36         4,39         2,50         4,39         2,50         4,39         2,50         4,39         2,50         4,39         2,50         4,39         2,50         4,39         2,50         4,39         2,50         4,39         2,50         4,39         2,50         4,39         2,50         4,39         2,50         4,39         2,50         4,39         2,50         4,39         2,50         4,39         2,50         4,39         2,50         4,39         2,50         4,30         2,50	Note	a		peggenn		30	4.15	•	3	97.9	6.39	<u> 5</u> .	203		•
ACCOLAGE         Inspectived         1,41         6,20         4,27         5,17         5,18         4,27         3,18           ALODOZEZ         District         Land         4,42         19,15         4         5,19         5,19         4,27         3,15         4,22         3,18         4,29         2,20         4,18         4,29         2,20         4,29         2,20         4,29         2,20         4,29         2,20         4,29         2,20         4,29         2,20         4,29         2,20         4,29         2,20         4,29         2,20         4,29         2,20         4,29         2,20         4,29         2,20         4,29         2,20         4,29         2,20         4,29         2,20         4,29         2,20         4,29         2,20         4,29         2,20         4,29         2,20         4,29         2,20         4,20         2,20         2,20         2,20         2,20         2,20         2,20         2,20         2,20         2,20         2,20         2,20         2,20         2,20         2,20         2,20         2,20         2,20         2,20         2,20         2,20         2,20         2,20         2,20         2,20         2,20         2,20 <td>AUCONOTION         Institute of the control of th</td> <td></td> <td></td> <td>. •</td> <td></td> <td>0.47</td> <td>2.07</td> <td>•</td> <td>ţ ;</td> <td>643</td> <td>17.7</td> <td>55</td> <td>3.88</td> <td>200</td> <td>•</td>	AUCONOTION         Institute of the control of th			. •		0.47	2.07	•	ţ ;	643	17.7	55	3.88	200	•
AUXILLAGE         Disast         V.A. 4.42         18.15         4.22         5.12         4.37         3.18           H1226.4         binast         Y         15.01         82.10         4         5.19         4.37         3.18           AAATJORD         binast         Y         15.01         82.10         4         5.91         4.39         2.50           AAATJORD         binast         Y         15.01         4         1.97         3.25         4.59           AAATJORD         binast         Y         15.01         4         5.81         4.59         2.50           H1368         binast         VAGO126         A.TZ         7.01         4         5.81         4.06         3.53           AAATJORO         binast         Link         6.81         4         5.86         5.86         5.89         3.23         3.50         4.06         4.06         4.06         4.06         4.06         4.06         4.06         4.06         4.06         4.06         4.06         4.06         4.06         4.06         4.06         4.06         4.06         4.06         4.06         4.06         4.06         4.06         4.06         4.06         4.06	H22854   Draft   F   F   F   F   F   F   F   F   F			pegipedami		1.41	979	<b>4</b>	7.77	2	4.23	10.05	2.98	1.07	, ,
ALTZBAS         briefled         192         6.25         4.5         5.19         4.5         5.5         4.5         5.5         4.5         5.5         4.5         5.5         4.5         5.5         4.5         5.5         4.5         4.5         5.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5         4.5	Mathods of the color	: :			¥	4.42	19.15	*	3.25		3.15	17.6	1.81	<del>1</del> 3.	•
Autocot         Desired         Y         15.01         02.10         4.58         5.48         5.58         4.59           HeBBAD         Coolen         Marybe         0.75         3.09         4         1.97         3.35         4.59           Auk43000         bone         y         1.77         26.06         4         286         2.48         4.59           W60136         sin         Marybe         6.37         26.06         4         3.88         2.48         4.08           Av70046         sin         13.56         54.17         4         6.81         4         4.08           Av70046         breast         VA         17.72         7.01         4         5.89         2.48         4.08           Av200126         breast         LA         6.81         4         5.19         6.11         5.02         3.89           Av200127         breast         col         4.18         6.41         6.11         6.02         3.01           Av200128         breast         col         4.18         2.44         6.11         6.02         3.01           Av200129         breast         col         4.16         4.16         4.17	Available         breast         Y         15.01         0.10         4.54         3.54         4.59         7.43         6.19           Hobbsolo         Color         Available         0.75         3.06         4         5.13         5.81         2.50         1.75           Available         sin         Maryon         0.77         0.10         4         5.89         2.89         4.50         4.28           Volume         vinit         Maryon         0.77         7.01         4         2.89         2.89         4.00         2.77         8.00           Volume         V         1.72         7.01         4         5.89         2.48         4.00         2.77         8.00         8.77         8.00         8.77         8.00         8.77         8.00         8.77         8.00         8.77         8.00         8.77         8.00         8.77         8.00         8.77         8.00         8.77         8.00         8.77         8.00         8.77         8.00         8.77         8.00         8.77         8.00         8.77         8.00         8.77         8.00         8.77         8.00         8.77         8.00         8.77         8.00         8.70 <t< td=""><td>• .</td><td></td><td>, je</td><td>i i</td><td>3.</td><td>8.25</td><td>•</td><td>5.18</td><td></td><td>92.6</td><td>3.26</td><td>4.67</td><td>2.90</td><td>•</td></t<>	• .		, je	i i	3.	8.25	•	5.18		92.6	3.26	4.67	2.90	•
AMAZINGA         Disease         Maybe         0.75         3.08         4         (97         3.33         7.00           AA443000         bone         Occion         Maybe         6.37         26.06         4         2.86         6.89         3.20         5.81           VMO128         sin         Maybe         6.37         26.06         4         2.86         2.48         4.06           H1308         bone         Y         172         7.01         4         3.89         3.21         2.71           VMO128         bone         Y         174         6.81         4         6.11         6.02         4.06           AA30276         break         1.74         6.81         4         6.11         6.02         3.34         1.74           AA30277         break         1.75         6.81         4         6.11         6.02         4.77         8.19           AA30772         break         1.20         4.18         6.84         4.47         8.19         4.77         8.19           AA30772         break         1.20         4.18         6.10         4.27         4.14         8.19           AA30772         break	MATINITION         Different         Matrix         Matrix <th< td=""><td></td><td></td><td>5</td><td>&gt;</td><td>15.01</td><td>62.10</td><td>•</td><td>5.48</td><td>9 0</td><td></td><td>7.43</td><td>6.18</td><td>126</td><td>•</td></th<>			5	>	15.01	62.10	•	5.48	9 0		7.43	6.18	126	•
Helbsto	AAAGSTOS         Cooler         AAAGSTOS         AAAAGSTOS         AAAGSTOS         AAAGSTOS         AAAAGSTOS         AAAAGSTOS </td <td>•</td> <td></td> <td>is read</td> <td>- 4</td> <td>0.75</td> <td>3.08</td> <td>•</td> <td>1.07</td> <td>S. S.</td> <td>3 6</td> <td>25.</td> <td>1.75</td> <td>4,44</td> <td>•</td>	•		is read	- 4	0.75	3.08	•	1.07	S. S.	3 6	25.	1.75	4,44	•
AAALOGO         Done         V/Met too         4.26.06         4.28.0         6.588         3.52.3           H13088         boses         Y         1.72         7.01         4         3.68         2.46         4.06           WOO128         bread         Y         1.72         7.01         4         3.89         3.21         2.71           WOO128         bread         N/A         1.44         6.81         4         6.17         5.05         4.00         3.94           AACORTOGO         hypochialemus         N/A         1.74         6.84         4         6.17         3.61         3.96           AACORTOGO         hypochialemus         N/A         1.74         6.84         4         6.17         3.62         3.89           AACORTOGO         hypochialemus         N/A         1.67         4         6.17         3.61         4.77         3.89           AACORTOGO         hypochialemus         N/A         1.67         4         6.10         6.17         3.89         4.77         4.77         4.77         4.77         4.77         4.77         4.77         4.77         4.77         4.77         4.77         4.77         4.77         4.77	AAA10000         Done         AAA10000         Done         AAA10000         AAA10000         AAA10000         AAA10000         AAA10000         AAA10000         AAA10000         AAA10000         AAA10000         AAA100000         AAA100000         AAA100000         AAA100000         AAA100000         AAA100000         AAA100000         AAA100000         AAA100000         AAA1000000         AAA100000         AAA1000000         AAA100000         AAA100000         AAA1000000         AAA100000         AAA1000000         AAA10000000         AAA1000000         AAA1000000         AAA10000000         AAA10000000         AAA10000000         AAA10000000         AAA1000000000	_		8	Š.	000	0.10	•	6.87	3.20	19.6	2.7	4.28	70.2	4
VR6189         stifn         Millsyca         24.72         7.01         4.06         2.48         4.06           H13088         bone         Y         1.72         7.01         4         5.89         3.21         2.71           MV00128         brasel         N/A         1.44         6.81         4         6.11         6.02         3.71         3.89           AA70046         hypothalanus         N/A         1.74         6.84         4         6.11         6.02         3.79           AA30277         brain         N/A         1.67         4.76         4         6.10         4.77           AA20670         bypothalanus         N/A         1.67         6.94         4         6.10         6.02         4.77           AA30277         brain         0.43         1.67         4         6.10         6.14         3.61         4.47         6.19         6.14         3.61         4.47         6.19         6.14         6.10         4.47         6.19         6.14         6.10         4.47         6.19         6.14         6.10         4.14         6.19         6.14         6.10         1.44         6.19         6.14         6.10         1.44         <	Wighting         skin         Million	•		pop ·			26.06	•	2.86	69.	3.5	150	80.8	3.94	•
H13088 bone   1156 5472	H13089 bons V 114 114 115 115 116 117 116 117 116 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 116 117 117	•		rjt S	Mayor	3 5	5	•	3.68	2.48	90.		90	320	•
WR0126         breast         134         589         144         681         4         611         632         389           AA702046         Indicated being	VR0128         bread         1,30         1,30         1,30         2,30         2,74         2,87           A/V00046         N/A (20046)         1,14         6,81         4         6,11         6,02         3,89         2,74         2,87           A/V00040         N/A (2014)         1,14         6,94         4         6,11         6,02         3,89         2,74         2,87           A/V00040         N/A (2014)         1,67         4         6,14         6,19         6,14         137         4,41           A/V00010         N/A (2017)         1,67         4,75         4         6,19         6,17         3,61         2,10         6,17         4,41         137         4,41         138         4,41         138         4,41         138         4,41         138         4,41         138         1,28         1,28         1,28         1,28         1,28         1,28         1,28         1,28         1,28         1,28         1,28         1,28         1,28         1,28         1,28         1,28         1,28         1,28         1,28         1,28         1,28         1,28         1,28         1,28         1,28         1,28         1,28         1,28         1,28 <td>YR.</td> <td></td> <td>Poog</td> <td>&gt;-</td> <td>* * *</td> <td>54.72</td> <td>•</td> <td>3.89</td> <td>321</td> <td>2.71</td> <td>707</td> <td>3</td> <td>231</td> <td>•</td>	YR.		Poog	>-	* * *	54.72	•	3.89	321	2.71	707	3	231	•
AATGO446         IVA         IV	AA70046         IAA	•		bre a A		867	1 1	•	1.27	3.01	3.86		70.0	3.41	•
Avazoros         hypochialismus         174         8 ms         4         2.08         6.00         4.77           Avazoros         brain         NA         0.61         1.28         4         2.08         6.00         4.77           R78247         brain         0.43         1.67         4         9.05         3.34         1.74           NAZZIZ         t-call         0.45         1.67         4         5.19         6.14         3.81           AAZZIZ         t-call         0.00         0.19         4         2.57         4.47         6.19           NAZZIZ         t-call         0.00         0.19         4         2.67         4.47         6.19           AAZZIZ         t-call         0.25         1.07         4         2.63         4.77         6.19           NAZZIZ         t-call         0.60         0.19         4         2.63         3.78         3.78           NAZZIZ         t-call         0.60         0.10         4         2.24         3.53         3.78           AAGSIGLA         bone         Y         0.86         3.26         4         4.17         3.18           AAAZIZES         provider <td>Avadestide         hypotinaliamus         174         0.54         2.08         6.00         4.77         5.89         0.11           Avadestide         hypotinaliamus         174         0.34         1.44         1.37         4.41           Avadestide         brain         0.43         1.67         4         5.19         6.14         3.81         2.10         3.38           NAZZIZ         t-cold         1.22         4.75         4         5.19         6.14         5.81         2.10         3.38         4.50           NAZZIZ         t-cold         0.26         0.19         4         6.40         3.83         3.79         6.53         7.21           NAZZIZ         t-cold         0.26         0.19         4         6.40         3.83         3.79         6.50         7.01           NAZZIZ         t-cold         0.26         0.10         4         6.40         3.83         3.79         6.50         7.01           NAZZIZ         t-cold         1.21         4.77         3.23         3.24         4.70         3.25         2.44         4.70           NAZZIZ         t-cold         1.21         4.67         6.41         3.23         <th< td=""><td>=</td><td></td><td>•</td><td>¥</td><td>=</td><td>i d</td><td>• •</td><td>6.11</td><td>6.02</td><td>3.60</td><td>2.74</td><td>Š</td><td>1.75</td><td>+</td></th<></td>	Avadestide         hypotinaliamus         174         0.54         2.08         6.00         4.77         5.89         0.11           Avadestide         hypotinaliamus         174         0.34         1.44         1.37         4.41           Avadestide         brain         0.43         1.67         4         5.19         6.14         3.81         2.10         3.38           NAZZIZ         t-cold         1.22         4.75         4         5.19         6.14         5.81         2.10         3.38         4.50           NAZZIZ         t-cold         0.26         0.19         4         6.40         3.83         3.79         6.53         7.21           NAZZIZ         t-cold         0.26         0.19         4         6.40         3.83         3.79         6.50         7.01           NAZZIZ         t-cold         0.26         0.10         4         6.40         3.83         3.79         6.50         7.01           NAZZIZ         t-cold         1.21         4.77         3.23         3.24         4.70         3.25         2.44         4.70           NAZZIZ         t-cold         1.21         4.67         6.41         3.23 <th< td=""><td>=</td><td></td><td>•</td><td>¥</td><td>=</td><td>i d</td><td>• •</td><td>6.11</td><td>6.02</td><td>3.60</td><td>2.74</td><td>Š</td><td>1.75</td><td>+</td></th<>	=		•	¥	=	i d	• •	6.11	6.02	3.60	2.74	Š	1.75	+
AASS275         brain         N/A         0.61         2.36         1.44         1.44         1.44         1.44         1.44         1.44         1.44         1.44         1.44         1.44         1.44         3.81         1.44         3.81         1.44         3.81         3.81         3.81         3.81         3.81         3.81         3.81         3.81         3.81         3.81         3.81         3.81         3.81         3.81         3.81         3.81         3.81         3.81         3.81         3.81         3.82         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.79         3.71         3.81         3.71         3.81         3.71         3.81         3.71         3.81         3.71	AASS275         brain         NA         OMI         238         1.44         1.37         4.71           R78247         brain         O.43         1.67         4.75         4.77         5.19         5.14         3.81         2.10         3.38           VAZ2772         t-aal         0.03         4.75         4         5.19         6.14         3.81         2.10         3.38           AASS772         t-aal         0.02         0.19         4         0.40         3.83         3.79         8.39         4.50           AASS772         brain         0.02         0.19         4         0.40         3.83         3.79         3.89         4.50           AASS778         brain         0.02         0.42         4         0.40         3.83         2.44         4.10           AASS778         brain         0.00         0.42         4         0.29         4.19         4.20         3.24         4.74         3.89         4.00           AASS778         brain         0.00         0.40         0.40         0.40         0.40         3.53         0.44         4.19         0.26         4.74         3.99         4.74         3.90         4.74 <td>3</td> <td></td> <td>hypothalamus</td> <td></td> <td>7.</td> <td></td> <td>• •</td> <td>2.08</td> <td>9.00</td> <td>4.77</td> <td>3.60</td> <td>7.</td> <td>1</td> <td>•</td>	3		hypothalamus		7.		• •	2.08	9.00	4.77	3.60	7.	1	•
R78247         brein         0.43         1/87         4         5/19         6/14         3,81           WQZZ72         Hoad         1,22         4,78         4         2,67         4,47         6.19           A/281742         Hoad         0,06         0,19         4         2,67         4,47         6.19           NS278         Hoad         0,06         1,18         0,68         4         2,93         4,18         2,58           NS378         brain         0,63         3,42         4         6,67         3,58         2,58           AA011320         color         3,58         13.06         4         6,67         5,61         2,53           AA03104         bone         7         0,86         3,56         4         2,67         5,61         3,56           AA03104         bone         7         0,86         3,56         4         2,67         5,61         3,56           AA03104         bone         7         0,86         3,56         4         2,67         3,51           AA03104         bone         7         0,86         3,56         4         1,17         3,61           AA04506	RT8247   Bright   Cold   Col	3		riero Liento	≨	6		,	9.05	3,34	<u> </u>	131		284	*
WQZZ7Z         Heal         1.22         4.76         4.76         4.77         6.10           AQZ8174Z         Leal         0.05         0.19         4         0.40         3.63         3.79           NZ5683         -         0.26         0.107         4         0.40         3.63         3.79           NZ5683         -         1.81         0.66         4         2.93         4.18         2.68           NS378         brain         0.83         3.42         4         2.34         3.33         3.82           AAG31014         bread         Y         0.86         3.56         4         4.16         2.79         3.16         3.56           AAAZ1628         preadule         Y         0.86         3.56         4         4.16         2.79         3.11         3.81           AAAZ1628         preadule         10.82         3.60         4         4.16         3.17         3.81           AAAZ1628         preadule         10.82         3.54         4         4.16         3.11         3.81           AAAZ1628         preadule         10.32         3.54         4         4.16         3.11         3.81	WG2772         Lead         1,23         4,78         1,27         4,78         1,29         7,51           AA261742         Lead         0,06         0,19         4         2,57         4,77         6,19         3,29         7,51           N25653         Lead         0,05         1,07         4         6,40         3,53         3,79         6,65         1,28         7,50           N25653         Lead         0,02         1,07         4         6,40         3,53         3,79         6,65         3,50         4,60           NS3778         brain         0,28         1,42         4         2,93         3,14         4,10         3,24         4,10           AA630104         brain         Y         0,88         3,49         4         2,67         5,65         3,56         4,74         3,35           AA630104         brain         Y         0,88         3,59         4         4,16         2,79         3,69         4,17         4,30           AA430104         brain         Y         0,88         3,54         4         1,73         6,18         5,19         5,19         2,19           AA447029         brain <t< td=""><td>5</td><td></td><td>Comple</td><td></td><td>3</td><td>) i</td><td></td><td>91.5</td><td>6.74</td><td>3.61</td><td>2.10</td><td><b>7</b></td><td></td><td>•</td></t<>	5		Comple		3	) i		91.5	6.74	3.61	2.10	<b>7</b>		•
AA201742         Leal         0.00         0.19         4         0.10         3.83         5.79           HZB683         -         0.28         1.07         4         0.40         3.83         5.78           HZB620         -         1.81         0.68         4         2.34         3.33         3.82           AA011320         colon         3.68         13.09         4         2.24         3.33         3.82           AA437162         bone         Y         0.88         3.36         4         2.22         5.15         3.36           AA437162         bone         Y         0.88         3.50         4         4.16         2.70         3.81           AA437162         bone         Y         0.88         3.54         4         1.73         6.29         3.11           AA437162         bone         Warm         11,33         40.56         4         1.21         4.61         5.01           TR2046         bone         Warm         11,33         40.56         4         5.81         3.15           TR2042         bone         13.21         4.61         5.01         3.17         3.80	AA281742         Leas         0.00         0.19         4         5.40         5.83         5.79         0.65         1.28           N28653	2		J.		ā	6.4		2.67	4.47	5.18	3.20	10.2		4
NESSES         0.28         1.07         4.18         2.68           NESSTR         brain         0.68         4         2.34         3.33         3.82           AAG01320         colon         3.68         13.06         4         6.67         6.61         2.65           AAA30104         bone         Y         0.98         3.36         4         6.17         5.61         3.36           AAA37528         preside         O.63         3.00         4         4.16         2.79         3.61           AAA37528         preside         0.63         3.53         4         3.55         6.47         3.81           AAA47628         preside         0.68         3.54         4         1.73         6.29         3.11           AAA47628         bread         II.33         40.56         4         1.21         4.61         5.01           T90990         bose         U.A         4         6.81         3.02         3.15           TR2842         bose         L.A         6.81         3.02         3.16         3.02         3.16           TR2842         bose         U.A         6.81         3.02         3.17         3.89	NGD663         LOT         LOT<	2				8		, ,	07.9	3.63	3.78	\$8,6	178	3 5	-
NSS78         brain         0.68         4.57         5.81         5.82           AA011320         colon         3.68         13.06         4.67         5.61         2.55           AA63104         bose         Y         0.88         3.56         4.16         2.79         3.56           AA421628         bosest         Y         0.88         3.50         4.16         2.79         3.56           AA421628         prosisio         0.63         3.53         4.16         2.79         3.81           AA421628         prosisio         0.68         3.54         4         1.73         6.29         3.11           AA441628         prosisio         0.08         3.54         4         1.73         6.29         3.11           AA441629         bosest         N/A         11,33         40.56         4         1.21         4.61         5.01           T90900         boses         N/A         11,33         40.56         4         5.81         3.02         3.15           T82642         boses         3.00         4         5.81         3.02         3.15           TR2642         boses         3.01         4         5.81	NS378         brain         0.68         4.73         3.33         3.82         2.44         4.10           AA011320         colon         3.42         4.67         5.61         2.65         4.74         3.65           AA011320         colon         3.68         13.06         4.67         6.61         2.65         4.74         3.65           AA03104         bone         Y         0.98         3.49         4.16         2.10         3.65         6.77         3.81         6.30         3.32           AAA37628         break         Y         0.98         3.54         4         1.73         6.29         3.11         1.20         3.32           AAA37629         break         N/A         11.33         40.56         4         1.73         6.29         3.11         1.20         3.12           AAA37629         break         N/A         11.33         40.56         4         1.21         4.61         5.01         4.50         4.57           TTD980         bone         N/A         11.33         40.56         4         1.21         4.61         5.01         4.50         4.57           TT234.2         bone         N/A         1.3	る		•		0.28	1.07	•	201	4.18	2.58	3.36	9.4	3 3	•
N83378         brain         0.89         3.42         4.87         6.61         2.65           AA011320         colon         y         0.88         3.36         4         4.67         6.61         2.65         3.36           AA427620         bose of one         y         0.88         3.45         4         4.16         2.79         3.62         4.77         3.81           AA427628         problete         0.88         3.54         4         1.73         6.29         3.11           AA045709         Neuron         VA         11.33         40.58         4         1.21         4.61         5.01           T90980         bose         VA         11.33         40.58         4         6.81         3.02         3.15           T82642         bose         VA         11.33         40.58         4         6.81         3.02         3.16           T82642         bose         VA         11.33         40.58         4         6.81         3.02         3.16	N83378         brain         0.60         3.42         4         6.61         2.65         4.74         3.85           AA011320         colon         3.66         13.06         4         4.67         6.61         3.55         6.77         3.85         6.77         3.85           AA03104         bone         Y         0.86         3.50         4         4.16         2.79         3.62         6.77         3.81         6.36         3.32           AAA37628         breast         10.52         38.08         4         4.16         2.79         3.61         4.50         2.38           AAA47628         breast         N/A         11.33         40.56         4         1.73         6.29         3.11         1.20         3.38           TT0080         bone         N/A         11.33         40.56         4         1.21         4.61         6.01         4.80         4.57           TT28-2         bone         3.30         4         2.09         3.17         3.80         1.92         6.82           AAA64104         N/A         0.30         0.30         4         2.09         3.17         3.80         1.92         6.82	20		•		1.0	8	• •	7.	3.33	3.82	7,7	<b>4.</b> 40		• •
AA03104         bose         Y         0.88         3.56         4         2.92         5.15         3.36           AA030104         bose         Y         0.88         3.56         4         2.92         5.15         3.56           HBZ081         bread         0.83         3.03         4         1.13         6.29         3.11           AA427528         proside         0.28         3.54         4         1.13         6.29         3.11           AA045709         Neuron         N/A         11.33         40.56         4         1.21         4.61         5.01           T9090         bose         N/A         11.33         40.56         4         5.81         3.02         3.15           T82842         bread         0.03         0.10         4         5.81         3.02         3.15	AA011320         Colon         3.66         13.06         2.22         5.16         3.36         6.17         4.30           AA030104         bone         Y         0.88         3.56         4         2.62         5.16         3.56         6.00         3.32           HR2081         breast         10.82         3.60         4         1.6         2.79         3.6         6.00         3.32           AAA27628         breaklike         10.82         3.80         4         1.73         6.29         3.11         1.20         3.28           AA045709         breaklike         NA         11.33         40.56         4         1.21         4.61         5.01         4.80         4.57           TR28-2         breaklike         NA         0.03         0.10         4         6.81         3.02         3.10         2.78         2.78           AA044104         NA         0.06         3.33         4         2.09         3.17         3.80         1.92         6.82	2		Ş		8	7	, -	197	5.61	2.55	4.74	3,85		• ◀
AA430104 bone Y 0.86 3.56 4.16 2.79 3.62 4.16 2.79 3.62 4.16 2.79 3.62 4.16 2.79 3.62 4.16 2.79 3.62 4.17 3.81 4.173 6.29 3.11 4.61 5.01 4.089 5.04 7.21 4.61 5.01 7.0090 bone N/A 11,33 4,056 4 1.21 4.61 5.01 7.02 3.16 7.02 3.17 3.80	AAA20104 bose Y 0.98 3.50 4 4.16 2.79 3.62 5.00 3.52 4.00 5.52 4.00 5.52 5.00 5.52 5.00 5.52 5.00 5.52 5.00 5.52 5.00 5.52 5.00 5.52 5.00 5.52 5.00 5.52 5.00 5.52 5.00 5.52 5.00 5.00	=		uojos		302	13.06	•	200	5.15	3.36	5.17	8	3 2	- ◀
H82061 bread 0.83 3.03 4 3.19 3.89 4.3.89 4.47 3.81 4.61 4.61 4.61 6.29 3.11 4.61 6.29 3.11 4.61 6.01 7.0090 bose N/A 11.33 40.56 4 1.21 4.61 5.01 7.02 3.15 7.02 3.15 7.02 3.15 7.02 3.15 7.02 3.17 3.80	Hazoet breast 0.83 3.03 4 3.55 6.47 3.81 6.38 2.39  AAA27628 preside 0.68 3.54 4 1.73 6.29 3.11 1.20 3.28  AAA477628 Neuron 0.68 3.54 4 1.21 4.61 5.01 4.80 4.57  Troppe bone NA 11.33 40.58 4 1.21 4.61 5.01 2.19 2.78  Trabat breast 0.03 0.10 4 6.81 3.02 3.15 2.19 2.78  AAA64104 NA 0.98 3.38 4 2.09 3.77 3.80 1.92 6.82	ö		powe	>	98.0	3.56	•		2.79	3.62	90.0	332	58.7	, -
AAA27628 proside 10.62 38.08 4 3.55 6.29 3.11 AAA427629 Neuron 0.28 3.54 4 1.73 6.29 3.11 TUQQQQ bose NA 11.33 40.68 4 1.21 4.61 5.01 TR2842 break 0.03 0.10 4 5.81 3.02 3.16	AAA27628 prosists 10.52 38.08 4 3.59 3.11 1.20 3.28 AAA27628 prosists 0.88 3.54 4 1.73 6.29 3.11 1.20 3.28 AAA427629 Neuron N.A 11.33 40.58 4 1.21 4.61 5.01 4.80 4.57 T00990 bone N.A 11.33 40.58 4 1.21 4.61 5.01 2.19 2.78 T72942 bread 0.03 0.10 4 5.81 3.02 3.15 2.19 2.78 AAA64104 AAA64104 AAA64104 AAA64104 AAA64104 AAA64104 AAA64104 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA	: 3		press		0.83	3.83	•	-	678	3.81	5.38	239	1.1	, ,
AAA45709 Neuron 0.28 3.54 4 1.73 0.28 5.01 T00980 bose NA 11.33 40.68 4 1.21 4.61 5.01 T02842 best 0.03 0.10 4 5.81 3.02 3.15 T32842 best 0.03 0.10 4 5.81 3.09 3.17 3.80	AAA45708 Neuron 0.88 3.54 4 1.73 6.28 5.01 4.80 4.57 70080 bose NA 11.33 40.58 4 1.21 4.61 5.01 2.19 2.78 782842 besset 0.03 0.10 4 5.81 3.02 3.17 3.80 1.92 6.82 AAA64104 1.00 0.00 3.30 4 2.00 3.77 3.80 1.92 6.82	:		- Indiana		10.52	36.08	•	CG.C		-	27	3.29	3,94	•
T00900 bose NA 11,33 40,69 4 1,21 4,61 0.01 T00900 bose 0.03 0.10 4 6,81 3.02 3.16 T02842 best 0.03 0.10 4 2.09 3.77 3.80	700000 boss NA 11,33 40,68 4 1,21 4,61 5.78  T00000 boss NA 0,00 1,02 3,16 2,19 2,78  702042 besset 0,03 0,10 4 6,81 3,02 3,16 1,92 6,82  AA864104 NA 0,96 3,39 4 2,09 3,77 3,80 1,92	3 :				8870	3,5	•	1.73	270		98.	4.57	128	•
T92542 break 0.00 4 5.81 3.02 3.10 T92542 break 0.00 4 2.09 3.77 3.86	1700800	2			41.4		40.58	•	121	4.61	9	•	2.78	4.53	•
T82842 bread U.S. 217 3.86	T02942 bread U.M. 0.96 3.33 4 2.09 3.77 3.86 1.84 U.M. AA664104	2	T90990		Š		Ç	<b>~</b>	5.81	3.02	3.10		3	3.86	•
	AA884104 - NA 0.98 5.30 .	2	T82842	Desag		3	2 6	•	2.09	1.17	3.86	1.82	•		
AA664104 - N/A 0.96 5.30		8	AA864104	,	Š	8	S								

Table 2.2: T3NO vs. BPH: Gene upregulated 5 fold or higher in at least 2 tumor samples or 3 fold or higher in at least 3 tumor Samples.

940	IMAGE	Gen Bank Accession	Tissue	Secretad/Me mbrane	AveBPH	AveTumor	FOIGIANETUMOITA FOIGINPMIZUA FOIGINPMIZIA FOIGINPMIZZA FOIGINPMIZZA FOIGINPMIZZA & OFTUMORS VERPH) V	Fold(MPM120/A	Fold(MPM121/A	FOLD(MPM124/A	Fold(MPM122/A	Fold(MPM123/A F veBPH)	-old(MPM12&A veBPH)	Samples
	5	Number		Bound							į	,	;	•
£	71011	44454004		N/A	298	10.39	•	3,92	5.10	3.87	3.78	7.7	4 .	, -
3				į	2	1.25	r	5.38	3.6	3.53	2.17	242	<b>9</b>	•
ž		2007	•	:	3 3			4.	3.12	4.85	17.7	3.40	6.29	₹
22	230	AA704650		>	5 5	8 5		2.26	5.31	3.00	<b>9</b> 0' <b>7</b>	4.15	2.00	•
128	89568	AA767401	<b>P</b> 000		2	8 :	, <b>.</b>	77.1	4.61	3,80	3.23	4.16	2.95	•
121	134507	P63263		>	=	90.	, ,	18.6	9	97	1.60	8.	3.28	•
128	289071	N63609		<b>&gt;</b>	38.8	8.5			<b>7</b>	13	4.28	8.4	38.1	•
Ę.	<b>40</b> 0203	AA677461		<b>§</b>	S :		, <b>.</b>	4.43	67.9	343	3.25	0.82	53.0	•
3	242081	185335	teada		27.	17.6	· ·		14	ğ	1.28	4.65	2.52	•
ē	20,08	H4502	erathyo		9.10	3 3	· •	187	5.62	3.24	98.4	1.60	3.25	•
132	306639	AA758154	color		8	224	, ,			204	98	8	3.08	•
<u>\$</u>	20608	H64086	ze Joseph	>	0.97	3.25	n .	2.04	3.48	i i	3.20	141	3.10	•
호	72897	ROBORA	٠		<b>3</b>	124	r	26.1	- (- 	8		707	20.	•
3	1467906	AABEDSO4	Prymeus	¥	1.18	7.0	r	1.87	3.62	3	)rr			
3	1656626	AA835580	elejeoud	*	1.17	3.77		3.58	9.6 9.6	320	<b>2</b>	3	<u> </u>	, ,
} ;	401/44		}	į	750	1.72	•	1.82	3.32	348	BC 7	3,33	3	•
2 :		2				12 24	•	1.60	3.05	3.27	4.18	6.11	<b>8</b> .	*
3	5	AA8Z4CS	2	Š	ò	47.0	• •	2.78	7	3,44	3.80	3.55	78.	•
2	282110	NESKY			, i	10.70	, ,		2.54	3.73	2.80	3.22	321	•
<del>2</del>	782047	AA429602	eye;eoxd	Maybe	1.1	3.6	, ,	07.7		9	3.87	3.10	4.59	•
7	454669	AA677208	•		ā	3.18 8.18	r	<b>N</b> 0.7	2 :	3 7		8	1.87	•
142	384283	AA709370		¥	9. 9.	3.	e.	4.76	7.00	<b>S</b>		į	2	•
2	240	R44132	Ç POR		0.53	3.1	n	2.04	<b>Z</b>	3.79	3.14		! !	, ,
: 3	743362	AAAOOSAA	eteleoro		8	000	n	3,33	3.10	2.05	4.16	98.0		•
	428785	810010	2	>	114	3,44	n	2.11	2.01	3.08	3.17	350	3.40	•
2 3		9		•	1.28	363		3.22	3.29	1.89	3,35	77	I,	•
2 ;		20000				131	r	3,15	2.85	325	3.12	38.	3.23	-
Ì						****		98.6	27.9	3.12	ā	0.47	3.55	<b>~</b>
<u> </u>		AA700804			3 :		, -	3.57	4.62	3.83	1.81	0.36	3.8	•
2	700	A4466348	5		= 6	3 8	. 7	2	1.13	1.13	**	134.44	1.13	n
<del>2</del>	710017	A12251		:	5 8	3 ;	3 %	2	1.72	0.11	0.61	3.63	22.02	
<del>2</del>	18.72	AA431721	e mana oud	-	3 3	2 6	2 ;		376	70.0	71.38	3.78	2.68	n
<u> </u>	220405	W1 8650	Poor -		8 8	8 5	<u>:</u> :	2	1.62	4.77	2647	8.4	202	n
3	2000	7				20.0	2 (		25.25	5.01	222	0.00	121	n
<u> </u>		1	endothelle		8 6		<u>.</u> t	7, 87	447	3	0.27	18.31	0.82	n
3		AA115761	entiple		\$ 1	5 5 6	<b>:</b> ;		0.0	2.23	65.00	5.81	3.4	c
<u>\$</u> !	19/04/	A443147	9000	;	3 6		= =	20.5	87 17	7.51	1.26	0.86	1.67	•
è		200	•	-	5 5	2 3	: Ş	2	97 07	4.51	28.1	101	228	n
3	200	148250	prostate		3 5	<b>1</b>	2 -	787	8	181	1.28	0.78	127	n
2		AA416627			3 3			1.07	47.4	18	5.30	32.20	2.64	n
<b>Ş</b>	470216	AA028987			2 !	78.7	• •		27.6	3	<b>3</b>	97.0g	2,42	n
<b>5</b>	376427	AA039851	7		ord ord	0.57	• (		7	3 5	12.30	20.5	0.12	•
ĝ	31	H49677	•		5	<b>1</b>	-	è i	2 ;	2 5		9	87	•
Ē	757451	AA437241	•	ΥN	0.47	3.37	-	4.67	<b>7</b> 8	0.7	8 5			
\$	466035	AA040461	musch	¥	8.25	35.89	-	2.41	677	2.60	10.17	900	3	,

Table 2-2: T3NO vs. BPH: Gene upregulated 5 fold or higher in at least 2 tumor samples or 3 fold or higher in at least 3 tumor Samples.

	#0414	Gen Bank	Tasue	Secreted/Me	Non-	AveTumor	Fold(AveTumor/A	Fold(Avelumoria Fold(MPM121/A Fold(MPM121/A Fold(MPM122/A Fold(MPM122/A Fold(MPM122/A Fold(MPM121/A	COLD MPM121/A	rold(MPM124/A veBPH)	Fold(MPM122/A veBPh)	Fold(MPM123/A	veBPH)	Samples
Orde	Clone	Accession	Distribution	Bound		Č	Vebru			į	5	147	324	•
		Nember			8	870	•	13.15	59.67	<u>r.</u>	8		180	n
3	298417	N74131	color			1	•	181	8,	<del>\$</del> 7	R of	977		•
į	666218	AAZ33609	rip(s		3	27	• •	; <b>;</b>	27	59'0	4.76	<b>9</b>	80.77	• •
! ;	087.480	CA177142	•		g	0.1	•	2 1	: :	2.16	10.05	<b>6</b> 07	*	,
è :		-	ļ		020	3,32	•	<u>.</u>	3 5	7	133	1.05	2.26	n
2	8	Variable V			0.81	4.56	•	5.88	10.01	; ;	220	980	2.12	n
<u>ş</u>	687687	WZ99370	hypornamics		8	32.96	•	6.81	14.24	20.5		17.07	0.10	n
570	712950	A282273	<b>Ž</b> .		8	0.43	<b>5</b> 0	6.26	15.0	7.72	2 6	9	2.62	n
17	504761	AA148735	Pove		3			3.13	90'4	6.73	20.30	3 3		*7
ţ	322440	W16424	adrenal		SI O	4		27.6	28,4	2.60	<del>3</del> .	11.48	900	•
	\$4010	81187	•		8	i.	•		3.74	8.08	86.0	7.48	878	,
3	2000	76747777	o adia		1.72	9.60	40	797	2 5	2.61	00.0	1.13	6.84	n
*				>	4	7.7	•	7.68	71.77		12.13	2,68	3.62	n
175	2012	20/62			27.0	60'17	•	1.82	8	<b>5</b>	2 6	7.72	4.30	n
176	11208	AA425650	rki.			90.0	50	1.52	1,24	13.33		į	98.0	•
Ħ	364865	AA035745	Dosest		\$ ;			98.1	14.29	<b>1.9</b>	<b>3</b> .	<b>L</b>		•
171	247567	NS8130	<b>Poot</b>		500	8	•		8	6.74	<u>.</u> 8	36.	3	•
	34888	R19878			914	0.67	n	9 9	7	4.47	1.57	4.55	0.13	•
<u>.</u>		144810	appearing		80	95.0	vo.	0.67			13.00	2,45	35.5	n
<u>\$</u>	3	200			12.74	56.08	\$	1.47	<del>-</del>	10.0	9, 4	1.05	7.64	n
<b>₹</b>	120	AA425682			20	90.0	•	1.67	6.21	2.89 2.89		96.0	77	n
182	11121	N57483	U BLA			5	ur.	7.28	8.45	6.19	ţ,	3 3	20	n
<u>5</u>	710088	AA430545	had		2	3		8:	6.33	7.29	3.	<del>2</del> ;		•1
Ì	61620	H19448	•		3	4 6		8.62	9.80	4.53	0.84	<b>S</b>	3 3	
2	256343	N26556	•		0.70	5.5		80 7	12.80	6.34	0.91	8:	<b>7</b>	• •
2	206170	W02426	•		S. C.	1.8/	•		1.86	1,80	5.23	124	3	•
:	778504	AM6139	prain		8	0.22	•	9 6		86.0	9.74	<b>3</b>	9.67	<b>.</b>
	73,404	AA128407	Drawal d		000	0.07	•	06.6 1	9,0	, c	2.13	9.89	<b>:</b> ;	n
3		227374	5015		90.0	0.28	₹	10.35	2 5		27.5	15.21	3.80	n
48 8	070870	Disease of	2		900	0.22	7	0.46	<b>8</b> .	86		101	1.74	n
<u>\$</u>	20702	Nester	•		(40	1.81	4	7.32	9.76	3.67	91.7	£	6.79	n
5	296907	N73563	•			7.18	•	7.89	2.40	5.57	97		570	n
ž	205628	H67585	prostate		5 6	2 8	•	5.29	7.00	99.0	<b>0.6</b> 0	8 .		e
5	67397	749326	ies is		00	3 5	• •	3.65	0.81	2,68	7.56	1.3	70.0	
ī	811028	AA485373	bressl		3	20.42	• •	,	98.	7.47	10.8	3. 3.	9C.1	
ā	4.89637	AA101878	prostate	ž	2.88	11.62	•			940	777	<b>3</b> .	. 22	•
	284624	N64796	•		1.56	6.38	•	A).7	2 2	4.10	3.36	1.15	208	•
:	841507	AA487385	or a	>	\$	4.28	•	10.80		0.28	9.36	0.79	X.	n
	A42407	AA678804	a de		900	0.14	*	0.28	3 8	1 .	2.63	4.80	90,4	63
2 :		107043			90:0	0.25	•	2.52	7.79		28.2	1,68	177	n
3	200	***************************************	,	<b>A</b>	94.0	1,79	4	2.83	1	970		1.83	11.30	n
ğ	1400/80	AA883/33	. !		100	0.10	*	3.06	<del>2</del> .	Ž,	8 ;	7	9.74	•
ž	784589	AA443300	<b>S</b>		3	80	•	3.91	7.78	1.48	0.70		91.4	**
202	723072	AA292683	pouq		ā 1	3	•	0.41	6.16	95.0	0.35	3	2 5	
g	656174	AA630620			27.0	5	•	55.8	7.03	2.57	0.65	0.88		
ğ	782575	AAA7522	COLVÁX		90.0	8. I	, ,	101	3.10	4.57	10.12	09.	1.70	<b>)</b> (
202	428186	AA001870	pooq		2.28	9.77	•		3 4 5	2.16	8.17	0; <del>4</del>	3.75	,
ķ	596626	AA130857	Neuron	₹ Z	1.17	4.49	*	2.78	2	•				
}								5						

Table 2-2: T3NO vs. BPH: Gene upregulated 5 fold or higher in at least 2 tumor samples or 3 fold or higher in at least 3 tumor Samples.

2.48 1.69 1.69 3.86 4.24 4.24 1.07 1.07 4.91 4.20 2.20 4.20 2.78 2.78 2.78 2.78 3.77
2.57 2.57 1.02 2.14 2.14 2.20 2.20 2.20
3.14 2.27 2.70 1.92 3.44 2.14 2.20 2.20 2.20 2.20 2.20 2.20 2.20 2.2
2.57 2.70 1.92 3.44 2.14 2.20 2.20
7.46 4.08 3.88
241 3.03 3.04 4.04
3,000
4 4 0.0
90'8
2.56 9.08
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250678 H95976 Ling Li

Table 2-2: T3NO vs. BPH: Gene upregulated 5 fold or higher in at least 2 tumor samples or 3 fold or higher in at least 3 tumor Samples.

Table 2-2: T3NO vs. BPH: Gene upregulated 5 fold or higher in at least 2 tumor samples or 3 fold or higher in at least 3 tumor Samples.

2.88 2.89 2.80 3.10 3.10 3.10 3.10 3.10 3.10 3.10 3.1					700	A LAND					17084	STORES CHORN CHORN	CHORN	Samples
WOODLITY         List         1156         3.15         3.15         3.17         3.18         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19         3.19	5	Accession	Olseribetion	Bound			veBPH)	<b>FAB9</b> 2	*****	A SELECT				•
March   Marc	QL W	******	di 4		8	3.15	ຄ	1.72	3.77	3.38	380	3	8 :	- 1
Marche   March   Mar	3 3	10000			27.00	75.03	r	3.30	3.07	2.81	8.	2.63	0.82	'n
Averages of season states         Av	3 3	- T	;			1	· e	3	321	1.05	29'9	1.07	4.77	•
AVINDADAS         Profit of the control of the co	3	AA461300	<b>3</b>	;	;	3 3		4.82	3.25	2.81	326	1.62	273	•
AMOTORS Name and AMOTOR	ğ :	AABBB10	prostele	¥ :	1.0	3 3	, ,	. S.	1.11	2.12	10.	3.10	1.04	•
AMSTATION         NAME AND COLUMN         NAME AND COLUMN         NAME AND COLUMN         135         135         136         136         136         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146         146 <td>9</td> <td>AA7023338</td> <td>Ę</td> <td><b>\</b></td> <td>2 3</td> <td></td> <td>, ,</td> <td>2.82</td> <td>3.27</td> <td>4.50</td> <td>8:</td> <td>1.83</td> <td>3.19</td> <td>n</td>	9	AA7023338	Ę	<b>\</b>	2 3		, ,	2.82	3.27	4.50	8:	1.83	3.19	n
Header	3	74877887	•	ž	<u> </u>	3	, -	1.35	3.30	3,86	2.74	3.60	2.47	n
March   Marc	3	8			7	3 2	ı e	5.57	3.34	17.1	234	0.80	701	•
Mail	ì	AA86344			2 2	3		2.76	3.94	2.31	3.06	2.04	3.43	n
Mail	3 3	128			3 5			3.82	7.58	3.41	1,10	1.06	970	n
March   Marc	3 3		٠		9.0	3		98	3.14	0.48	0.21	77	22	n
ALAMENTOS         Decided of the control of the c	3 3	A4001	. }		3 5	<u> </u>		2.20	263	3.17	2.07	3.85	<u> </u>	-
Maintain			į		3 8			7.	3.16	1,41	5.6	4.41	<u> </u>	•
Accounts         Prof.	2 8	A4450			3 5	40.0	, en	87.7	2.11	3.29	157	3.43	8,	n
	3 :	AA660/ 60		<b>\frac{1}{2}</b> :		5		308	3.4	3.69	2.10	ភ	1.7	n
M51468         brish         0.01         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03	3	2002	prostate	<b>&gt;</b> -	•	3 5		180	377	9,1	123	8.	g	n
Mid-148   Drain   V   O.45   L.200   D.24	3	AA062813	<b>e</b>		5	3 3	, .	5	4.75	3.79	3.	0.30	36	•
W/70147         V         0.40         1.31         3         2.01         3.24         3.18         3.52         2.33           R84243         benna         V         0.40         0.13         3         2.59         4.53         3.04         2.48         3.30         2.33         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30         3.30 <td>3</td> <td>N51496</td> <td>e la la</td> <td></td> <td><b>6</b>.</td> <td>8</td> <td>n (</td> <td>3 :</td> <td>191</td> <td>95.9</td> <td>1.78</td> <td>121</td> <td>1.74</td> <td>•</td>	3	N51496	e la la		<b>6</b> .	8	n (	3 :	191	95.9	1.78	121	1.74	•
House   Marcol   Ma	8	W70147		> :	<b>9</b> 9	2 ;	, c	2 6	3.24	87.13	3.52	233	2.40	
NAJ2522         busine         γ         1.69         4.01         3.89         3.24         2.59         3.41         1.52           AAAB0133         busine         γ         1.69         4.01         3         3.73         4.87         3.45         1.24         1.80           AAAB0731         busine         γ         1.51         1.55         3         2.39         4.27         3.40         1.24         1.80           AAAB0731         busine         0.02         1.14         3.59         3         2.39         3.43         1.24         1.80           AAAB0722         brain         0.02         0.14         3         0.24         0.29         3.44         1.80         3.44         1.80         3.44         3.84         4.45         3.84         4.45         3.84         4.45         3.84         4.45         3.84         4.45         3.84         4.45         3.84         4.45         3.84         4.45         3.84         4.45         3.84         4.45         3.84         4.45         3.84         4.45         3.84         4.45         3.84         4.45         3.84         4.45         3.84         3.84         3.84         3.84         3.8	3	H08725	prostate	>	2.7	- ·	· •	5	127	3,04	2.48	3.30	0.85	n
MASSES         National State         1 cm	8 :	X82475	<b>9</b>	:	5 5		, •	5	324	2.58	3.41	1.50	219	•
AMMENS ALL         CANADER ALL         1,31         3,59         6,23         3,40         3,07         2,53           AMMENS ALL         CAMPERS ALL         1,31         3,59         3,23         6,23         3,40         3,97         2,53         3,44         3,44         3,44         3,44         3,44         4,45         4,45         4,45         4,45         4,45         4,45         4,45         4,45         4,45         4,45         4,45         4,45         4,45         4,45         4,45         4,45         4,45         4,45         4,45         4,45         4,45         4,45         3,47         3,23         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22         3,22 <t< td=""><td>2 :</td><td>N.25282</td><td>5 2</td><td><b>&gt;</b></td><td>9 6</td><td>? .</td><td></td><td>2.5</td><td>197</td><td>3.45</td><td>124</td><td>1.80</td><td>2</td><td>c</td></t<>	2 :	N.25282	5 2	<b>&gt;</b>	9 6	? .		2.5	197	3.45	124	1.80	2	c
Automotion         Automot	<b>R</b> :	AMMOIS			3 -		s er	2.36	623	3,40	2.07	2.63	0.52	•
AAA45702         brain         0.445         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.54         0.52         0.54         0.51         0.52         0.54         0.51         0.52         0.54         0.57         0.52         0.54         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57         0.57	3 :				3 5	; ;		5.37	900	138	1.18	3.6	3.62	•
Availation         Paris         Color	9 :	18/8074	. :		7 6			3	920	330	2.89	57.4	4,47	n
No. 1553   No. 1554   N. No.		AAA > 102	ş			} =	, -	2.13	3.70	2.31	1.13	3.56	3.32	n
NST 553 Unitoperation         NA.A.D.4221 placemia         NA.A.D.4222 placemia         NA.A.D.4	3 5	HORUGE		1	3 6		. ~	191	3.40	1.91	3.28	322	272	n
MAXILIARY         Presents         2.18         5.81         3.99         3.90         3.23         2.04         1.07           MAZZBA         AMAZZBA         Co.4         0.11         3         1.59         3.47         3.00         0.99         2.09           AMAZZBA         AMAZZBA         O.64         0.11         3         0.21         4.19         1.79         4.62         0.21           MAZZBA         NAM         0.87         2.28         3         0.21         4.19         1.79         4.62         0.21           AAAZBAS         Namera         NAA         0.87         2.28         3         4.47         0.62         1.83         3.29         0.70           AAAZBAS         Namera         0.87         2.27         3         3.24         1.65         3.29         3.72           AAAZBAS         Namera         0.87         2.27         3         3.21         3.07         3.61         2.20         3.72           AAAZBAS         Namera         0.87         1.14         3.02         3.01         1.14         3.20         3.01         3.20         3.20         3.20           AAAZBAS         Namera         0.80	÷ ;	28.58	peupedaun	ž	, R	7 6		2	231	1.16	348	3.11	4.17	n
NAZORA         NAZORA         O.O.H.         0.11         3         1.59         3.47         3.00         0.09         2.00           ANAZORA         NAZORA         O.O.H.         0.11         3         0.21         4.19         1.79         4.62         0.21           HOSORO         NAZORA         O.O.H.         0.13         3         0.21         4.19         1.79         4.62         0.21           AAZORA         Nacora         O.O.H.         0.10         3         0.20         3.01         2.54         1.65         3.20           AAZORA         Drumin         O.O.H.         1.70         3         4.47         0.62         1.89         3.70         1.62           AAZORA         Drumin         O.O.H.         1.71         3.02         3.21         3.07         3.61         2.30         1.62           AAZORA         Drumin         O.O.M.         1.17         3.02         3.71         3.01         3.72           AAZORA         Drumin         O.O.M.         1.54         3         2.00         2.31         3.20         3.21           AAZORA         Drumin         O.O.M.         0.04         3         0.02         2.31	;	15000	an and an			3		3.69	3.90	3.23	204	1.07	505	•
Available         Available         Available         6.021         4.18         1.78         4.62         0.21           H45206         H45206         0.06         0.13         3         0.21         4.18         1.78         4.62         0.21         3.33         3.58           H80696         Heart         0.87         2.28         3         4.47         0.62         1.87         3.28         3.28           Avaistage         Heart         0.87         2.27         3         3.21         3.07         3.61         2.30         1.62           Avaistage         Free Inc.         0.87         2.27         3         2.27         3.61         2.30         1.62           Avaistage         Free Inc.         0.87         2.27         3         2.00         2.01         1.14         3.90         3.01           H6744         Dearer         1.22         4.32         3.27         3.29         3.01         3.20         1.62           Avaistage         Free Inc.         0.90         1.54         3         2.00         2.01         3.20         3.21         3.20         3.21           Avaistage         Free Inc.         0.90         1.54		1000	Į.		2 2	1		95:1	3.47	3.00	80	2.90	3.07	•
H45289         NA         0.87         2.28         3         189         202         372         3.33         3.59           N80989         Neart         0.87         2.28         3         4.47         0.62         180         2.54         185         3.29           AAA28174         bread         0.87         2.27         3         2.27         3         2.54         185         3.29         0.70           AAA28174         bread         0.87         2.27         3         2.21         3.07         3.61         2.30         0.70           AAA28171         bread         0.87         1.77         3.02         3         1.62         3.07         3.59         3.01           H6744         bread         0.90         1.54         3         2.09         2.01         1.14         3.90         3.01           AAA88602         bread         0.04         3         0.02         3.70         2.31         0.04         3.72           AAA47126         bread         NA         6.57         1.66         3.02         3.24         1.79         4.79         4.79         4.44         0.58         1.77         4.79           AAA	3 5		•		9	61.0	· m	0.21	4.18	1.79	4.52	0.21	808	•
N85089         Near         178         3         203         3.01         2.54         1.65         3.29           AA480399         AA480399         Lide         4.39         3         4.47         0.62         1.83         3.69         0.70           AA291749         Dread         Lide         4.39         3         4.47         0.62         1.83         3.69         0.70           AA291740         Dread         Lide         2.27         3         3.21         3.07         3.61         2.30         1.62           AA291740         Dread         Lide         1.37         3.27         3.21         3.59         0.77         3.59         0.77           AA29174         Dread         Lide         1.34         3.07         2.01         1.14         3.90         3.01           H15744         Dread         Lide         3         0.62         3.70         2.31         0.64         3.72           AA498602         Lide         3         1.39         3.09         1.64         3.37         0.77         0.24           AA497126         Dread         1.00         1.00         1.00         1.00         1.00         1.00         1.	2 3	277			240	2.28	- 67	1.80	202	3,72	333	3.56	8	•
Audelists         Lies         4.39         3         4.47         0.62         1.83         3.99         0.70           Audelists         Dread         0.87         2.27         3         3.21         3.07         3.61         2.30         1.82           Audelist         Dread         U.A         1.17         3.02         3         1.18         1.22         4.32         0.72         3.59         1.07           TP6151         Naminal         0.90         1.54         3         2.09         2.01         1.14         3.90         3.01           HIST44         brain         0.02         0.04         3         0.62         3.70         2.31         0.64         3.72           Audelists         brain         0.02         0.04         3         0.05         3.00         1.84         3.72         3.41           Audelists         breat         0.02         0.06         3         1.09         3.09         1.04         0.32         3.41           Audelists         breat         0.02         0.06         3         1.37         0.44         0.58         1.17         4.50           Audelists         1.02         0.02         0	Ş	100000	. }	É	5	1.79		2.00	10.6	2.54	1.85	3.26	301	n
AA221740         bread         0.87         227         3 321         3.07         3.61         230         1.62           AA421740         bread         N/A         1.17         3.02         3         116         1.22         4.32         0.72         3.59           AA42151         Namon         0.90         1.54         3         2.09         2.01         1.14         3.90         3.01           H16744         brain         0.90         1.54         3         0.02         3.70         2.31         0.64         3.72           AA457126         breat         N/A         6.57         16.82         3         1.39         3.09         1.94         2.32         3.41           AA622432         N/A         0.72         0.06         3         1.37         0.24         3.09         4.79           AA622432         N/A         0.72         0.09         3         1.17         4.50			į		9	4.39	. 10	15.4	0.52	8.	3.00	0.70	81	n
Autotack         Mark         1,17         3,02         3         116         1,22         4,32         0,72         3,59           TP6151         Namon         0,90         1,54         3         2,09         2,01         1,14         3,90         3,01           H15746         brain         0,02         0,04         3         0,62         3,70         2,31         0,64         3,72           Autostato         NIA         6,57         16,82         3         1,39         3,09         1,64         2,32         3,41           Autostato         NIA         0,72         0,09         3         1,39         3,37         0,77         0,24           Autostato         NIA         0,74         1,89         3         1,17         4,50         4,79	3 5	97270474	j		0.17	127	. 67	3.21	3.07	3.61	2.30	1.62	<b>L</b> .	n
TP5151         Nauron         0.90         1.54         3         2.09         2.01         1.14         3.90         3.01           H16744         brain         0.02         0.04         3         0.62         3.70         2.31         0.64         3.72           AA488602         NIA         6.57         16.82         3         1.39         3.09         1.64         2.32         3.41           AA457126         heart         0.02         0.09         3         1.32         2.29         3.37         0.77         0.24           AA828432         NIA         0.74         1.89         3         1.17         4.50         4.79         4.44         0.58         1.17         4.50			į	**	1 12	3.02	n	1.16	121	4,2	0.72	3.58	29	n
H16744 brain 0.02 0.04 3 0.02 3.70 2.31 0.04 3.72 A448602 NJA 6.57 16.82 3 1.39 3.04 1.94 2.32 3.41 A4487126 breat 0.02 0.09 3 3.82 2.20 3.37 0.77 0.24 A482322 NJA 0.74 1.89 3 1.37 0.26 0.92 3.00 4.79 4.44 0.58 1.17 4.50	j	706.6	1	<u> </u>	90	4	n	2.00	201	1.14	3.80	3.01	131	<b>5</b>
Addition heart 10.02 0.09 3 1.39 3.09 1.94 2.32 3.41 Add37126 heart 0.02 0.09 3 3.82 2.20 3.37 0.77 0.24 Add27126 heart 0.04 1.89 3 1.17 0.26 0.92 3.00 4.79 Add2712 1.89 3 1.17 0.26 1.97 4.40 0.58 1.17 4.50	3				0 02	00	n	0.62	3.70	2.31	<b>3</b> 9.0	3.72	4.51	n
AAAST126 heer 0.02 0.09 3 3.82 2.20 3.37 0.77 0.24 AAB22432 NAA 0.74 1.89 3 1.17 0.96 0.92 3.00 4.79	3 4		Š.	W.W	15	16.82	. 63	1.30	906	7.	232	74	3.2	n
AAB22432 NAA 0.74 1.80 3 1.17 0.96 0.92 3.00 4.79	;	7	. ]	Š	2	5		3.82	2.20	3.37	0.77	0.24	<b>1</b>	n
A682432 NA 0.14 1.54 5.50 1.17 4.50	3	971/5	1	į	7 6	8 .	י ר		8	0.92	3.8	£.4	#	'n
	7 9	AA642452	•	Š	7 0	8 6	ກ ເ			\$ F	1.17	8:	0.30	-

Table 2-2. T3NO vs. BPH: Gene upregulated 5 fold or higher in at least 2 tumor samples or 3 fold or higher in at least 3 tumor Samples.

		Gen Bank	i	Secreted/Me			Fold(AveTumor/A	Fold(MPM120/A	FoldiaveTumoria Fold(MPM120A Fold(MPM121A Fold(MPM124A Fold(MPM122A Fold(MPM121A Fold(MPM125A # of Tumors)	Old(MPM124/A	Fold(MPM122/A	Fold(NPM123/A	Fold(MPM125/A	s of Tumors Samples
O S	IMAGE Class to	Accession	Distribution	mbrane	AveBPH	AveTumor	veBPH)	wBPH)	v•BPH)	VeBPH)				
		Number		Bound	;	900		3.31	3.85	3.58	1.37	1.10	3 ;	, ,
ä	503741	AA131486	brain		2	207	, ,	7.7	94	0.76	35.	3.28	67.1	2
ž	764130	AA432075	prostete	Maybe	17.73	1.11	י מ	5 6	7.87	2.82	1.13	0.31	314	n
2	325370	W52208	bone		9.03	90.0	en (	9.00	3	0.85	1.80	1.37	3.87	n
2	288663	N62394	Mdney		0.78	 2.	n	3.07		41.0	3.08	7.02	0.08	n
11	344359	W73377	gun		0.07	0.1 <b>6</b>	2	<b>9</b> 6.0	7	75 13	34.	123	1.35	•
	6330	R16241	٠.		1.15	2.79	~	3.60	6	1.87	201	1.08	3.55	n
} }	429823	AADOBBOS	•		0.7	0.35	2	3.16	- C	. c	4.12	0.85	232	n
}	258742	NYOOKS	•		0.62	1.40	2	3.61	3.63	3 5	1.52	3.40	3.14	n
} ;	758807	NOT YOU	•		000	0.20	~	0.60	<b>7</b>	3 5	97.6	3.01	1.30	n
į ;	800.37	44707436	trae 4		8	8.3	~	1.29	324	9 9		0.38	3.18	•
7 :	LACT TO	AAAAAAA	atalana		20.08	46.9	7	2.43	08.5	6.5	. S. E	0.24	4.05	•
? :	A028C1	731010	a property		900	6.0	~	1.20	3.87	<b>7</b> 0	2 2	99'0	28.0	n
;	BA00F4	AA487506	ş		0.36	0.76	7	4.12	3,45	7 6	72.0	42.46	32.68	~
3		44469492	and the line		5	0.18	13	0.74	0.74	8 6		27.75	5.82	2
<b>3</b> :	2000	2000	oler!		800	0.50	•	1.60	0.16	7.78 7.78	19.5		244	7
3	200	738CE	i :		92.0	2.25	•	8.43	28,00	0.37	800		¥.	7
3	137864	R63065	5		3	ğ	•	1,46	0.70	1.19	57.0	* ·	. 346	. ~
ž	854585	AA660222	)		4 5			7.32	50,05	8.	0,49	9 1	3 :	
ş	151477	H02837	akin			2 g	. ~	15.33	16.31	2.50	<u>4</u>	99.0	B **	• •
ž	1331	111091		>	<b>i</b>	2 ;	. *	13.67	22.88	0.57	3.	1.01	87.0	• •
355	82029	H22932	•		<b>Q</b>	8 5		1.07	0.12	0.12	0.52	31.38	97.5	
ş	129630	AA886405	nolos		8 8	3 ;	•	880	0.41	0.38	1.35	18.46	16.17	•
켥	99062	AA006196	•		ឌ	<b>3</b>		98	2,38	80.1	0.55	1.30	20.60	N9 (
35	204446	N21233	•		8	7	, .	2	7.7	1.77	2,77	11.30	1200	<b>N</b>
88	754358	AA438142	bone		2	3 .	•	240	191	16.0	20.25	2.06	5.30	~ ₁
35	757 165	AA443850	Neuron		8	0. 5	n •	48.4	14.43	2.52	1.47	0.61	2 <del>4</del> 0	N .
3	865238	AA138125	Deser de		18.25	80.08	n u	2 2	0.10	0.65	0.22	9.95	2,80	<b>7</b>
88	47355	H10981			0. 5	0.48	n ·	3 5	7,	11.68	178	230	2.82	7
98	377896	AA777098	16.834	Ş	9. 9.	16.28	₹ .	5.12	9	123	14.1	1.67	0.83	~
8	745077	AA626275		ž	3	1.83	•		40	1.07	1,48	800	79.71	м
382	70763	160081	fund		0.16	0.7	•	: :	10.86	0.50	0.62	0.21	1.48 84	<b>14</b>
2	1380018	AA855158	e logo		8	0.20	•	1 33	10.63	5.68	2.67	1.68	8	<b>19</b>
¥	283335	N5141			2	0.72	•	: #	8.36	680	0.57	0.82	10.57	η.
386	28483	N71861			20 0	0.0	•	20.7	0.36	6.13	7.	0.36	16.08	~
*	1031203	AA609982			900	ב ב	• •	: ;	9	2.86	27.7	8.	99.0	<b>N</b>
×	1048988	AA778636	endothellat	¥	0.14	0.57	•	2.5	200	2.96	8.45	<b>9</b> .00	7.	~
8	46160	AA682274	breest	¥	10.27	38.61	•	<b>1</b> 1			2.21	5.65	6.87	~
8	60692	H17626	Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda Punda		15.0	1.18	₹	1.87	187	3 5	16.11	17.0	0.71	~
5	347928	AA018980	Ģ		101	8	₹	2.78	776	3 2	3	070	10.61	*
; ;	177178	TORATE	•		500	2.0	*	0.95	5	*		0.52	33.0	7
5 5	20804	NEGOTA	٥		950	200	•	6.43	12.29	P. 1	8 6	6.30	1.99	~
•	101866	140360			0.61	77	•	2.33	8.78	2.53	8 6	610	R	7
	165072	27130			0,0	101	•	7.00	9.56	2.16	7.0	•	!	
Š								o						

HAGE	Gen Bank	5.2	Secretedilie	Aughbu	AveTumor	Ahor	FOICHMPM120/A	Fold(MPMM24A	FOIGIMPMIZUA FOIGIMPMIZOA FOIGIMPMIZUA FOIGIMPMIZUA FOIGIMPMIZUA FOIGIMPMIZUA FOIGIMPMIZOA FOIGI	Fold(UPU122/A	FOIG(MPM123/A veBPH)	FOR WERN	Samples
`_	in the second	Distribution	Bound			(Hage)				. !	٤	4.74	7
				;	70,	•	1.28	2.12	2.17	7.62	4	3	, ,
•	1134862	Pone		2.	i,	•			80	8	10.16	<b>8</b>	~
	57.963	Pees		о О	8	<b>~</b>	9.0	3 1	;		1.85	78.2	~
	063230	Pusch		88	0.10	•	6.17	150	8 7	2 5	989	2.39	2
	<b>670280</b>		ş	782	<b>8</b> .	n	1.42	1.16			961	1.86	7
	159620	bresst		98	1.82	n	2.20	\$ 70	1.01	47.6	0.15	69'0	~
	74257			0:02	<del>ب</del>	6	6.65	<b>6</b> .18	16.7	2	8	90'9	~
	V480115	9		9.12	39.44	n	1.38	0.0	98.	773	7.72	225	*
x	115366	•		770	0.71	n	0.69	9.		3	0.65	901	~
3	003028	•	¥¥	<del>5</del> .	18.4	n	5.71	7,10	87 :	3 5	98.5	1.00	*
₹	1022201	٠		7.	6.67	n	5.38	2.73	8 8	3 5	1.89	128	~
₹	1629024	teatte	¥	171	3.80	n	2.60	5	976	3	\$50	0.70	~
T	116743	brain		2.07	6.83	e	2.59	20.0	2 2		0.26	8.48	~
\$	977CBV	prostate		700	0.10	n	09:0	0.26	9.6	<u>.</u>	790	6.62	7
₹	W63518	gur,		0.13	0.32	7	0.33	0.07	· ·	8 6	25	5.32	7
_	78886	prostate		90	1.45	7	0.73	124	<b>9</b>	•			

Table 2-3: T3NO vs. BPH: Genes upregulated 10 fold or higher in at least 2 tumor samples or 5 fold or higher in at least 3 tumor samples.

	a sa nasduma.	tumor samples of e total or me					Aloca Montage	FoldIMPM121/A	FORGWPM124/A	Fold(MPM) Zon	VARIATION TOTAL FOLDING FOLDING FOLDING FOLDING TOTAL CONTINUE TOTAL VARIATION VARIATI		Samples
	Gen Bank	l	Secretedilite		A. Tumor	Fold(AveTumouAv	Halam.	veBPH)	veBPH)	(HARA)		,	•
			<b>Markine</b>	Avenu		2	•	į	4.28	62.32	12.54	800	•
	Clone ID Number	Parameter 1	Bound	;	14.544	67.98	149.63	166.70	1	31.06	62.59	8.8	•
78.6	DARKED AAAKTOID	operate of		8.		47.98	65.24	24.62		A 26	41.24	19.50	ь ,
5				2.38	14.00	7.00	31.27	20.03	À		45.42	20.45	•
į	•		>	2.41	E 5	7.56	20 80	11.57	48.9	/c.er		19,03	•
Ř	284701 N64840		-	2.85	78.64	30.0B	30.05	8	10.09	19.72	21.41	3 9	•
4	450049 AA703398	os prostate	ž	3	0.21	22.48	1,34	8	23.40	15.57	11.61	ATC	•
7.				100		17.62	22.19	20.62	3 1	10.00	5.71	721	•
ξ '				24.06	473.60	9 9	87.00	41.52	0.28	ñ :	\$0.24	8.	•
2	2810CS N.50880		>	2.56	44.38	3.5	7	7. 7	6.16	924	201	***	•
ţ	136605 R35051	-	-	•	25.46	17.23	30.48	<b>.</b>	8	10.01	17.00	8 1	•
ţ	171130 R28397	atalamq 7	<b>&gt;</b> -	<u>?</u>		11.81	9.28	\$. =		7 12	8.55	19.24	> 1
: :		e pou		0.0	•		6.78	6.32	80,00		12.71	17.9	•
			,	2.07	20.30		68 10	9.60		1 2	67 64	10.0	•
9				0.50	3,63	19.7		2.45	<b>18.9</b>	12.55		1.76	₩,
=======================================	196975 R91398			Ş	3,88	12.94	V2.CI		10.24	6.27	02.51		~
	287528 NG2096	. 2		3 5	13.52	10.47	7.81	24.30	10 70	41.0	4.27	3.5	•
	259374 N31952	52 prostate		9.		10.16	7.09	15.81	2 :	43 CF	2.63	8.0	
			>	124	12.01	4	5.69	10.94	8.72	<u>.</u>	5.75	6.20	•••
				0.27	201	3 5	3.24	10.05	6.80	9.7	16.8	11.57	*
15	•			0.20	1.97	29.0		2.	5.38	5.13		2.08	10
16 2	21931 772543		•		12.19	6.37	970		5.30	6.38	8	3 5	•
7	825859 AA504784	. 751	2		0.20	5.62	7.23	2 1	72 03	105.24	0.80	3	•
	130136 R20813			5	200	34.16	18.89	29.		980	23.62	128	•
			¥	3	3	22.20	29.78	48.11	007 <b>8</b> 2		15.40	1.72	•
		enod erry	ž	9. 9.	1230	7	9.65	9.81	<u>-</u> <del>3</del>	3 9	13	3.92	4
				0,40	7		680	28.16	14.54	17.70	! !	3.27	*
E				15.97	184.72	•		18.00	79.7	13.10	<u> </u>	2	7
# F	`		>	17.59	158.73	8.02	65	40.46	13.83	13.07	8.		7
2	193923 R83407	101 purp	-	2	0.0	9.80	13.49		7.85	4.72	3.65	170	• •
	467165 AA045074				2	8.70	13.54	10.00		10.01	4.07	6.1	•
	305086 N91003	86	>	2	270	6.73	3.84	10:11		10.48	20.7	263	•
	22217 N79180	180 · bone		8		6.55	7.58	10.38	70.0	200	ī	3.81	•
				0.10	77.		5.05	10.82	5.48	2	77.	10.42	•
			ž	0.38	2.39		88.8	5.08	10.11	A1.2		2.67	•
		G G G G G G G G G G G G G G G G G G G		11.01	70.00		8.5	*	5.12	20.9	E		4
29	1056172 AA62	AA620995 muscle		800	5.55	5.84	8.95		12.8	7.48	3.67	9.40	• •
9	762363 AAA31407					5.84	3.19	6.0	2 6	6.43	5.07	2.77	•
	195034 R88	R85764		70.		5.48	2.60	6.47		1.30	0.26	S.D.8	•
_		- v4706258	ž	X i			6.20	78.	8		3.0	0.17	•
: :		AAAS2278 prostate		1.76			6.74	6.67	0.33	1	5.02	1.7.1	•
3		A A 4.700KB		0.02			40.5	6.48	9.30 •	3	113	7.07	•
z				0.47	2.07			5028	3.85	18.7	40.0		•
3	195367 R&			3.44	46.10	13.39	8.		503	171	0.0	4 :	•
8	83888 AAK	AAAS7235 prostate	_			12.85	43.16	70		0.27	18.31	0.82	•
: 2	281039 NA	N47717 endotheltal	3	3		12.37	46.34	0.67	2 :	29.8	48.21	5,35	n
; 5	_	4A115761 embryo		\$			3.78	5.13	8				
				5	27.02	11.81	7.7	!					

Table 2.3: T3NO vs. BPH: Genes upregulated 10 fold or higher in at least 2 tumor samples or 5 fold or higher in at least 3 tumor samples.

		Gen Benk	i	SecretediMe			ENJARATIONALE FOLDIMPHIZIZE FOLGIMPHIZIA FOLGIMPHIZIA FOLGIMPHIZIA FOLGIMPHIZIA FOLGIMPHIZIA	Fold/MPM120/A	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A	FOID (MPM123/A		# of Tumor
500	Clone ID	Accession	Distribution	mbrane	AveBPH	AveTumor	•BPH]	₩8PH)	VeBPH)	\HdB•×	veBPh)	√eBPH)	(HAB)	
		Number		Bound	į	;	8	08 04	24.83	7.51	125	98'0	1.67	n
\$	827786	N60168		>	0.51	6.43	10.05	70.88	2 1		0.28	2.33	1.01	-
7	18473	AA779728	proclete	¥	0.13	1.36	10.30	22.40	13.74	3.4			2.78	•
: ;	7670571	4 4 847 997			800	0.69	8.20	10.16	27.83	<b>3</b>	63	\$ !		
; ;		********			60	0.11	8.02	10.01	67.0	2.20	24.30	4.57	5.0	•
2 :		MAN 1348	j		7.	11.25	6.59	7.30	5.16	4,03	15.83	2.66	4	, מ
:	8	PC7/-					9.48	8.59	3.86	13.77	2.12	<b>8</b> .38	<b>6.</b> 18	n
2	20021	N20066		:	ì	3 6	<b>3</b> 54	9.37	12.80	2,46	6.80	235	3.29	•
2	50103	AA165678		<b>-</b>		5 6	<b>.</b>	. A	3	9.60	4.70	4.07	3.73	•
2	428816	AA004681			2 :	8 5		0.12	3	17	<del>1</del> 00	2.60	8.45	n
3	11582	AA454587	popo	>	200	77.17		: :	<b>7 7</b>	4.73	5.71	11.11	7.68	n
9	426438	AAD04412	•		3	717	3.6	2 6	3	2	11.80	***	2.0	•
3	795729	AA440291	petpedin		200	 -:	100 S	8 1	76.75		0.72	99:0	2.12	•
2	712050	A282273	3		g	<b>8</b>	80	0 0		90.		10.6	4.50	n
3	825785	AAS05117	OV#Y		154	8.57	5.51	3.24	B :			17.071	010	•
3	504751	AA148735	poue		90.0	0. \$3.	6.45	6.26	re.p	777	2 6	8	2.67	•
3	481706	AA115259	Pre-sei		97.0	3.80	5.23	3.83	6.17	<b>5</b>	7000			
2	241621	H83381		ş	8	5.43	5.10	3.85	1979	5.92		27		, -
. 5	782537	********			1.72	9.60	5.01	2.52	2.75	80.0	90.0	₹ '	970	
2 2	277594	M43003	2	>	99.	7,	4.07	7.68	12.12	2.17	000	1,13	\$ 1	٠,
: :	10763	1000 B	1	. >	1.25	6.16	4.92	2.53	5.09	3.89	<del>*</del> .46	6.48	5.1	<b>.</b>
2 5	97770	20000	. }	•	0.20	04:1	4.85	5.02	7,58	4.08	<b>1</b> .0	3.53	2.03	n i
2 9			3 1		58.0	45.42	4.71	5.23	10.10	4.4	5.26	1.16	207	-
, .					0 0	0.0	6,70	0.48	4.83	6.38	7.51	<b>1.</b> 30	7.58	•
		9,0070	•		71.0	0 67	4.62	5.69	9.05	6.74	8.1	8.	2.03	n
	200	0/961X	. !		12.85	50.29	197	5.52	8.50	8.8	1.01	£.63	<b>9</b>	n
, ,		138/4	and di		0 0	800	8	1.67	173	2.55	1.18	1.05	. 7.08	•
•	904.00	701	5 5		5	15.28	3,	9.25	5.31	0.40	5.30	3.29	363	n
	#2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/00 #2/	27,000			3 5	33.38	4.52	7.28	8.45	8.19	7.0	0.28	2.38	n
	F18.20	2000	5		3	0.12	4.40	8	8,33	7.29	3.	2.43	<b>8</b> 6'0	n -
	136.907	20000			•	8.20	4.38	2.17	5.43	1.71	<b>X</b> :	3.80	5.05	n
	738604	00000	de		8	0.22	8,	0.98	7.86	1.80	5.23	1.24	<b>X</b>	-
	205628	S S S S S S S S S S S S S S S S S S S	or a fair		150	2.18	2	7.09	2.40	29'9	1.28	58.1	6.78	n
,	30000		1		8	0.20	27	0.16	3.28	3.22	<b>9</b> .00	8.0	6.10	n
- :	707	T40734	1		100	900	4.18	5.29	7.89	0.68	08:0	99.0	9.46	n
• •	7674		1		5	5	84	2.09	6.18	6.46	2.24	6.60	34.	-
2 3	107.00		. j		3	71.0	18	0.28	7.63	0.28	9.35	0.79	<b>3.</b>	•
	76107		į		100		88	6.41	<b>6</b> . 10	9670	0.35	133	6.10	•
2 9			j		0.12	70	3.78	29.1	000	1.78	6.80	8.74	151	n
2 :	01.00	987.			9	7.	3.78	5.28	285	X.7	1.60	9.76	1.01	n
: :	2000	38278244	and a		190	8	9.40	1.86	1.13	5.78	5.41	1.13	<b>8</b> .00	n
	177047	73067			100		25 92	7.0	1.13	1.13	14.34	134.4	1,13	~
2 9	1000	*162CA	1		3 3	3 5	18.02	69.03	<b>3</b> .5	3.22	1.15	1.7	15.6	~
Q :		78677987	and and	;	5 5	5 5	16.42	64.82	77	0.11	Q.61	3,63	20.22	~
=	22228	AA431721	prostate	<b>&gt;</b>	3	<b>4</b>	19.42	78:40	<u>‡</u>	: ;	į			

Table 2-3: T3NO vs. BPH: Genes upregulated 10 fold or higher in at least 2 tumor samples or 5 fold or higher in at least 3 tumor samples.

8 of Tumod Stamples 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2
FOLCHMISSIA VARPH) 2.05 32.88 22.8 4.21 1.40 0.38 16.87 12.03 1.46
Fold(upustable) veBPH) 44.85 42.49 1.07 0.78 1.04 1.04 1.14 11.20 0.21
Fold(MPM122/A veBPh) 28.47 0.74 1.88 1.28 1.64 1.63 2.85 2.85 2.77 0.52
FOLGIMPMIZUA ve BPH) 4.77 1.96 4.51 1.81 2.50 0.57 4.37 1.77 0.50
Fold(MPM1711A veBPH) 1.52 - 0.74 28.40 28.89 19.31 22.89 2.31 22.89 2.31 18.83 0.41 10.89
Fold(MPM120/A veBPH) 0.73 0.74 21.88 14.84 15.33 15.33 13.67 3.07 13.15 0.88 2.04
Post(aveTumordav         Fold(luPM120IA         Fold(luPM122IA         Fold(luPM12IA         Fold(luPM12IA)         Fold(luPM
AveTumor 12.90 0.18 10.34 33.07 3.19 1.66 5.97 0.48 0.48
AveBPH 0.96 0.01 1.90 4.08 0.47 0.05 0.05
SecretedMe Ribrare Bound Marbs
Tas ue Discributiba spieen spieen endothelisi prossti  colon bone
Gen Bank Accession Number Not111 AA180111 HZ2832 R39068 N74131 AA005196 AA38142 AA38142
BAAGE Clame ID 20053 20063 731280 81331 52028 731280 81331 52028 7418 742086 75438
O

Samples

Table 2-4: Poor vs. Good Clinical Outcome: Genes upregulated 3 fold or higher in any one tumor sample.

upregulated 3 fold or higher in any one tumor sample.

Table 2-4: Poor vs Good Clinical Outcome: Genes

Fold(AvsPoorIA Fold(mpm31/A Fold(mpm32/Av Fold(mpm33/A # of Tumors 62.14 10.17 7.73 6.80 3.28 3.60 0.86 0.33 8.85 2. 6.75 6.98 6.70 9.92 5.95 5.88 6.62 4.97 5.81 ¥. 0.67 12.28 6.19 3.83 0.7 0.03 0.75 2.47 6.00 0.70 2.50 0.66 0.26 0.00 0.36 1.37 5. 62.30 25.33 17.80 10.19 3.29 7.66 1.28 6.82 5.91 3.61 8.87 6.78 1.58 6.64 6.55 5.82 9.84 2.18 4.21 5.87 2.52 0.85 0.93 1.06 0.92 6 1.59 0.75 0.69 3.69 4.12 4.90 8 0.45 0.24 0.81 0.47 0.82 0.02 1.37 1.66 0.37 5.95 3.39 1.55 veGood) 31.26 5.49 3.95 3.83 5.61 4 4.23 3.52 3.51 3.47 3.28 3.19 3.17 3.17 2.97 2.67 9.81 4.25 3.81 5.17 5.0 AvePoor 169.35 108.65 52.25 19.88 1.50 237.74 16.09 7.78 21.23 2.09 0.16 3.83 12.41 2.53 3.89 8.54 6.95 3.63 2.37 1.47 5.16 3.29 6.32 AveGood 24.18 13.03 9.8 53.79 53.51 0.86 0.5 8 2.03 0.80 0.05 1.20 3.91 9.0 <del>...</del> 3 0.87 1.24 0.27 4.08 0.38 6.0 1.47 AA677708 AA700876 Accession AA436479 AA708152 AA630084 AA099262 **4A478298** AA488676 **AM32030 AA432030** AA001431 AA677388 W93315 W84751 AA028171 W79647 N76201 N58558 H68848 R10935 N92842 H91680 AA406301 R02586 AA664101 R61073 AA444051 IMAGE Clone ID 753076 248412 212188 782513 782513 129128 854678 382059 308696 415768 489535 364865 754559 756595 452374 357105 460798 460487 454822 241705 487327 124071 346321 855624 42325 287528 284601 740941 <del>1</del>6 ဖ 5 4 5 oder

Table 2-4: Poor vs. Good Clinical Outcome: Genes upregulated 3 fold or higher in any one tumor sample.

4.79 4.42 4.95 5.26 5.26 3.86 3.81 3.81 3.82 3.83 3.49 3.49 3.49 3.49 3.49 3.49 3.49 3.4
1.18 4.85 0.29 0.29 0.83 2.12 2.04 1.06 3.41 3.04 0.94 0.94 0.94 0.94 0.94 0.94 0.94 0
4.79 6.06 6.28 4.59 3.14 4.20 1.28 1.28 1.28 1.29 3.03 3.03 3.03 3.03 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16 3.16
0.99 1.32 0.83 0.62 1.13 1.19 1.19 1.11 1.13 1.140 1.140 0.57 0.94 1.28 0.08 0.08 0.55 0.56 0.56 0.56 0.56 0.57 0.66 0.72 0.74 0.75 0.75 0.75 0.75 0.75 0.75 0.75 0.75
2.94 2.90 2.90 2.90 2.50 2.50 2.34 2.34 2.35 2.35 2.34 2.34 2.34 2.34 2.35 2.35 2.34 2.34 2.35 2.36 2.37 2.38 2.39 2.30 2.30 2.30 2.30 2.30 2.30 2.30 2.30
4.06 4.08 1.86 0.02 43.42 1.70 3.00 19.45 15.91 0.19 8.05 7.96 69.88 4.15 0.13 1.05 1.05 1.05 1.05 1.05 1.05 1.05 1.05
0.22 1.48 1.10 0.67 0.01 18.45 0.05 1.20 1.17 1.17 1.19 1.20 2.08 2.08 2.08 1.81 2.09 0.21 2.09 0.21 2.09 0.21 2.09 0.25 0.25 0.25 0.26 0.27 0.05 0.05 0.05 0.05 0.05 0.05 0.05 0.0
N73958  AA456886  AA432049  W46900  AA682386  AA428415  W84716  AA428415  W84716  AA489743  AA4897499  W42812  AA489743  AA46174  H86554  AA461774  H86554  AA461774  H86554  AA461774  H86554  AA461774  H86554  AA461774  H86554  AA465888  N90051  N51486  AA460511  H25546
298303 NY 298303 NY 298303 NY 461759 AA 461759 AA 461759 AA 461759 AA 415692 NY 415692 AY 415692 AY 415692 AA 41214 F 898096 AV 430231 AA 430231 A 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099 303099
8 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5

Table 2-4: Poor vs. Good Clinical Outcome: Genes upregulated 3 fold or higher in any one tumor sample.

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12.08	0.42	1.00	14.50	13.33	13.47	0.62	1.33	11.93	1.09	1.48	13.64	0.61	2.41	8.86	0.48	0.84	0.20	9.83	0.73	8.89	1.16	0.79	0.45	1.30	1.20	1.27	0.51	6.89	1.71	1.55	1.10	2.15	2.30	2.18	
1.16	0.97	13.86	0.84	1.28	0.83	12.60	2.26	0.90	12.18	08.0	0.40	1.26	1.81	2.20	0.70	0.66	<b>1</b> .0	0.86	0.34	96.0	0.78	2.68	1.13	1.64	1.08	1.00	7.98	2.38	1.72	0.77	2.52	0.62	1.70	1.42	
1.61	0.92	0.70	1.15	0.94	0.90	0.52	10.78	1.21	1.09	1.48	0.31	11.31	1.97	1.14	12.18	0.92	0.96	0.92	0.73	2.00	9.88	1.70	1.47	1.30	1.20	1.81	0.57	0.90	7.08	2.28	2.89	2.16	2.01	1.41	
2.95	15.34	1.91	0.51	1.38	0.67	2.12	0.84	1.08	0.62	11.18	0.43	1.03	7.91	1.73	0.40	10.97	11.67	1.64	11.16	0.87	0.83	6.86	8.81	7.61	8.34	7.27	2.20	0.92	0.56	6.45	4.47	5.87	4.79	5.78	
4.45	4.41	4.37	4.25	4.23	3.97	3.97	3.80	3.78	3.74	3.73	3.69	3.55	3.53	3.48	3.44	3.35	3.32	3.31	3.24	3.18	3.16	3.01	2.97	2.96	2.96	2.84	2.81	2.77	2.77	2.76	2.75	2.70	2.70	2.70	,
0.82	0.39	3.58	2.10	4.89	4.59	6.59	2.51	3.06	3.20	5.85	18.28	0.32	73.02	3.01	0.21	0.73	0.15	2.94	8.46	0.85	1.53	48.46	6.88	13.27	2.40	0.93	5.99	3.09	4.88	0.09	6.25	32.44	2.47	2.89	
0.18	0.09	0.82	0.49	1.16	1.16	1.68	99.0	0.81	98.0	1.57	4.95	0.09	20.71	0.87	90.0	0.22	9. 8.	0.89	2.61	0.27	0.48	16.11	2.32	4.48	0.81	0.33	1.08	1.11	1.76	0.03	2.28	12.01	0.91	1.07	
AA782380	AA863204	RG3823	AA704718	W69950	H41222	H19128	AA628410	N48261	AA425382	W38679	AA451886	H06093	N64840	T77410	W47487	N33041	T99719	AA458413	W48762	H38425	H12981	AA136808	AA678880	N91797	N47738	H17882	H44861	W31255	R00480	AA180608	A133721	AA486281	AA598659	AA024898	
857442	1455740	141336	450948	343919	192258	50930	1032774	282117	73157	320857	786657	44351	284701	23790	324313	270343	122782	788386	325128	192401	43207	491001	1493160	307174	277423	50162	176904	306540	123331	592707	512116	842848	897901	365227	
65	99	49	68	69	20	77	72	23	74	75	92	22	78	79	8	₩	83	83	\$	82	88	87	88	88	8	9	35	93	94	92	98	26	88	66	

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6.67	0.71	47.4	<u> </u>	76.0	1.96	0.94	1.75	1.41	15.	2.93	2.01	1.15	2.36	1.80	6.80	3.38	2.70	0.26	0.55	1.81	1.33	1.39	1.49	2.10	0.72	69'0	1.24	6.99	1.59	6.65	2.40	1.29	0.89	1.26	0.54	;	
80	8.5	7 9	84.	96.0	2.34	0.92	2.75	1.08	1.45	4.64	1.56	77.0	223	0.69	1.41	2.86	2.51	0.50	£0.	28.4	1.45	3.61	2.19	2.30	2.03	1.21	2	0.72	1.60	960	188	4 92	1 83	58.C		<u>.</u>	
!	1.47	1.13	0.67	1.14	1.26	13	3, 1	2,25	7.5	7.13	7 7		 	181	50.0	2.54	3.23		0.90	· ·	05. ±	97.6	2. C.	0 i	. e	4.32	57.5	100	<u> </u>		2 5	0.32	<u> </u>	D 10	0.63	[ <del>-</del>	
	1.67	7.75	6.78	1.92	483	9 6	n (	00.4	0.40 0.60	80.4	1.46	5.73	9.90	3.52	0.00	0 0	1.13	0°.1	8.20	6.74	1.20	9.00 0.00	2.68	2.28	3.35	0.80	6.32	3.80	0.70	0. 13 10. 13	0.61	4.5g	1.88	1.96	6.89	6.08	
	2.69	2.68	2.67	2 66	90:	7.90	2.60	2.56	2.55	2.54	2.54	2.52	2.50	2.50	2.49	2.48	2.47	2.45	2.45	2.45	2.44	244	2.43	2.43	2.41	2.39	2.39	2.37	2.37	2.36	2.35	2.34	2.34	2.34	2.33	2:32	•
	2:38	2.19	2.51	4	5.73	21.09	0.73	4.25	7.65	3.12	7.59	14.37	13.42	0.07	34.17	1.57	2.60	5.92	23.38	6.48	7.65	0.13	2.62	3.96	6.17	39.	4.56	99.4	2.42	0.38	1.74	30.32	26.12	0.54	0.13	6.68	
	0.88	0.82	9	5	<del>-</del>	8.12	0.28	<b>3</b> .	3.00	1.23	2.39	5.71	5.37	0.03	13.72	0.63	1.05	2.41	9.53	2.65	3.13	0.05	1.08	1.63	2.56	69.0	1.91	1.96	1.02	0.16	0.74	12.93	11.15	0.23	90.0	2.88	
	AA701931	AA885311	110000	007670	N51367	85797N	H75860	AA682631	R17982	W75603	R92601		AA010932	AA676470	AA443004	R15409	T95274	AA865464	AA621031	AA629991	AA434362	AA704844	AA278408	AA778098	AA421389	AA999953	N27982	N76101	AA628146	R12183	AA757732	R98064	R11587	AA621324	W02342	N26163	
	135567	433301	#010#t	35	283208	289589	230560	431296	32381	345670	196257	361363	359781	882511	811766	53162	120162	1470048	1056203	884514	837908	453218	703930	379920	739237	1636495	265102	299412	1055737	25590	396111	206838	130046	1048794	292071	269411	
	5	3 5	5 5	<b>1</b> 02	103	\$	105	106	107	108	109	110	=======================================	112	113	114	115	116	117	118	13	120	121	122	123	124	125	126	127	128	129	130	131	132	133	\$	

-		-	-	-	<b>-</b>	-	-	-		- •	- ,	- •	-	-	-	-	-	-	-	-	-	-	-	-		-	-	-	-	-	-	-	-	-	-	
1.28	0. <b>83</b>	0.78	2.08	0.52	0.76	0.43	1.85	92.0	2 4	9 k	c	3.42	1.52	0.51	0.80	1.86	1.66	0.81	1.55	0.53	3.05	5.49	1.70	5.74	1.18	1.09	1.09	4.10	2.24	1.13	0.95	2.18	0.92	3.17	0.97	
2.50	1.79	1.52	2.27	1.21	5 15	- F.	5 5		1.40 0.10	8/7	1.20	1.63	2.50	0.85	0. <b>8</b> .0	1.71	3.14	1.03	<u>.</u> 4	<del>.</del> 4	2.50	1.15	1.33	<u>2</u> .	<del>2</del> .	1.52	2.08	1.37	0.92	0.92	0.90	<del>2</del> .	5.68	1.66	5.76	
0.87	2.68	1.74	3.23	1.62	<u> </u>	27.	11.7	00.1	0.75	3.29	1.80	2.82	1.36	1.13	6.45	1.57	1.13	1.02	1.67	1.92	2.59	0.86	0.67	1.13	1.55	2.34	2.19	1.13	2.24	1.13	1.29	1.52	1.29	1,19	0.97	
4.59	3.93	5.17	1.59	08.4	3 6	20.7	3. A.	2.29	6.15	1.56	4.74	1.21	3.67	6.43	0.88	3.81	2.99	6.07	4.18	4.93	0.68	1.28	5.06	0.80	4.62	3.71	3.29	2.04	3.19	5.41	5.44	3.03	79'0	2.51	0.82	
2.31	2.30	2.30	333	67.7	67.7	2.29	2.29	2.28	2.28	2.28	72.2	2.27	. 2.26	2.26	2.24	2.24	2.23	2.23	2.21	2.21	2.20	2.20	2.19	2.18	2.17	2.17	2.16	2.16	2.15	2.14	2.14	2.14	2, 14	2.13	2.13	4
6 62	23.40	4.7.4	t :	<b>5</b> (	0.13	4.17	19.01	1.33	0.75	4.37	4.23	1.08	85.0	1.52	đ	2.12	41.6	0.76	9 6	10.94	80	1.43	7.68	50.0	98:	8.	28.92	5.24	8.31	90.65	27	5.69	0.58	11.18	1 14	,
4 20	200	0.70	0 (	3.90	8	1.82	8.32	0.58	0.33	1.82	1.86	970	0.43	990	670	, yo	3 7	<b>7</b>	5 6	3 6	4.60	3. 6.	3 5	200	, Y	0.83	13.37	2.43	3.87	28.28	0.57	2.66	0.27	5.24	3	;
1008004	1000704	AA19/344	AA8/3088	W76331	AA444111	AA676899	AA011583	H15250	R44776	AA733090	N45398	AA406059	ACAZON.	HAKABO	000000	A & CORESCO	A5577368	44824457	A A 0.4754 B	A071844	900000	N63630	A4142813	4402424	A A D D 4 9 7 3	87898N	AA171899	AA134862	AA454165	AA011095	AA419612	H53141	AA777417	AA251386	A A 6 70 4 29	!
307000	285403	627688	1325751	345055	757135	455179	429678	49687	33086	398577	260187	743026	30000	10000	10501	240400	08/80	1044747	717001	20070	1364303	C70001	17767	364473	478078	28274	61m74	AU2478	795322	140881	748148	202521	140061	AB4687	200 <b>-</b> 00	
, ,	200	38	137	138	139	140	141	142	143	144	145	146	7 5	<u> </u>	9 9	1 4 4 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5	5 5	55	727	3	<u> </u>	22	00.	) q	2 4	2 6	2 2	£ 5	4 5	3 5	ž	9 4	3 5	2 4	00 9	2

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4.99	4.82	0.84	1.32	1.89	4.53	1.65	3.11	2.79	1.39	0.25	3.92	<del>7.</del>	1.78	0.62	5.97	0.15	1.50	1.36	0.84	1.12	1.74	1.34	2.51	94	0. 44	1.67	0.59	69.0	2.01	0.65	0.65	0.68	1.39	0.62	
1.09	1.69	1.07	4.86	0.92	1.88	1.47	1.24	3.06	1.24	2.62	1,57	0.96	3.50	1.09	0.71	0.13	1.68	3.03	0.70	2.17	1.67	1.57	0.12	3.10	0.99	1.23	£.	1.49	0.99	1.52	1.72	1,53	4.20	1.31	
,	ر د د	8 8	200	5 +	8 5	2 2	£ 5	1 16	136	3.23	0.86	6.73	1.79	1.05	09'0	7.72	1.29	1.36	5.44	1.12	1.74	4.37	5,34	2.82	2.51	4.23	1.63	1.81	2.01	1.97	3.14	2.06	1.19	22.1	
•	1.07	1.07	5.25	09.1	3.70	68.0	y	ξ, γ,	1.31		 	. 54 44	1 2 1	5.47	F 6	9 6	0.20	2.12	4 K	5 6	3.04 4.04	2 6		2 72	4 18	65 C	55.	4 11	- 60 - 6	75	75.0	3.79	4 28	4.88	9
	2.12	2.12	2,11	2.11	2.10	2.10	5.09	2.08	2.08	2.07	2.07	2.07	2.07	2.06	2.06	2.05	2.05	2.05	2.5 4.5 5.6	2,04	2.04	2.04	2.03 2.03	2.03	2.63 50.5	2.03 503	37.7	2.03	2.02	7.07 7.03	2.02	2.02	2.02	2.02	10:7
	0.78	80.65	3.29	13.63	13.41	3.21	2.34	27.88	2.66	0.79	36.21	0.69	1.92	3.48	2.14	16.88	9.61	6.73	1.09	0.27	2.06	3.34	0.75	18.02	13.69	4.76	0.77	5.26	4.26	1.38	26.52	9.71	7.47	3.17	0.24
	0.37	42.83	1.56	6.45	6.39	1.53	1.12	13.37	1.28	0.38	17.47	0.33	0.93	1.69	<u>5</u>	8.23	4.69	3.29	0.53	0.13	1.01	1.62	0.37	8.88	8.75	2.35	0.38	2.60	2.11	0.68	13.11	4. 48.	3.71	1.57	0.12
	AA706955	W96187	H83283	AA626242	AA010223	AA521414	N47308	AA455565	AA453040	AA676948	AA876375	N71769	AA187789	AA030046	W58013	R45964	. AA031513	AA011185	AA430242	95660H	T61792	AA676840	AA872602	AA151002	AA968896	N20328	AA598572	AA609983	AA164782	AA150507	W48700	H67292	AA490985	R10948	H09749
	464808	401090	400475	745524	430235	926090	280508	813488	788574	460189	1493107	290702	624379.	469969	341336	35575	470393	359713	781091	46916	78946	460114	1323328	504927	1674594	264597	897822	1031978	595001	491763	324885	229573	824588	129148	46453
	ļ	5 5		2 <u>.</u> 2 .	5 5	174	5 5	177	178	179	180	181	182	183	\$	185	186	187	188	189	190	191	192	193	194	195	196	197	198	199	200	201	202	203	204

<b>-</b>	_	-	-	-	-	-	-	<del>-</del>	-		<b>-</b>	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	<del>-</del>	<del>-</del> -	-	-
£.1	1.24	1.02	08:0	0.79	0.59	3.62	1.46	1.1	0.80	3.20	1.30	0.75	0.51	1.50	1.78	1.31	57.0	1.30	6.57	2.06	0.82	1.01	0.97	0.93	3.07	<b>.</b> 08	0.89	1.10	0.93	1.65	1.13	0.97	2.39	0.0 0.0
1.35	2.50	1.20	1.69	1.29	1.17	1.20	4.84	3.49	1.22	1.25	1.07	1.82	1.38	1.09	1.28	3.99	2.79	3.96	0.42	2.03	1.05	0.93	1.03	1.19	2.59	1.15	0.76	1.00	1.26	3.06	4.27	1.19	3.27	0.67
1.73	0.76	0.96	1.88	4.31	5.08	1.38	0.80	2.01	1.69	4.4	1.55	0.76	0.95	1.33	1.54	1.11	3.06	1.63	0.40	3.64	3.99	1.02	0.98	0.94	0.82	1.10	2.06	2.00	1.74	1.80	1.14	1.71	1.18	6.77
3.60	3.49	4.80	3.49	1.56	1.09	1.71	0.81	1.29	4.19	2.02	3.98	4.53	5.03	3.92	3.22	1.40	1.22	0.87	0.36	0.00	1.86	4.75	4.71	4.62	1.20	4.33	3.95	3.55	3.70	1.10	1.07	3.66	0.75	0.05
2.00	2.00	1.99	1.99	1.99	1.98	1.98	1.98	1.98	1.98	1.98	1.97	1.97	1.97	1.96	1.96	1.95	1.95	1.94	1.94	1.93	1.93	1.93	1.92	1.92	1.92	1.92	1.91	1.91	1.91	1.90	1.90	1.90	1.90	1.89
5.87	9.35	0.91	19.17	2.63	0.16	10.70	3.81	06.4	8.22	3.66	0.25	20.0	0.09	2.75	171	1.89	33.80	1.30	151	0.03	4.71	<b>3</b> 50	10.1	0.41	4.14	0.68	8.	55.28	6.25	1.81	0.56	<b>8</b> .	3.29	4.03
2.93	4.68	0.45	9.63	1.32	0.08	5.41	1.93	2.48	1 16	1.85	0.13	9	40.0	1.40	0.87	260	47.34	0.67	0.78	0.0	2.44	0.28	0.63	0.21	2.16	0.35	1.03	28.91	3.28	0.95	0.29	4.71	1.73	2.13
T83657	N51362	R31114	AA195601	R07948	W02073	N26008	AA879435	AA664105	W15319	N62293	AA455825		W32729	AA259115	AA169173	H38263	AA399074	A A 309433	AAAJAASA	H04828	AA634379	70077	H29290	R39364	AA449048	H20717	R01245	AA453105	AA478606	H03146	W79069	R92801	R23609	171061
113284	283191	134265	665127	127063	327247	268837	1505688	855834	122643	20024	811609	64272	327495	686684	810006	190950	729953	775800	770340	43968	743828	23218	49858	24632	785816	51420	124137	789091	753683	151597	346968	197056	132140	84078
205	206	202	208	506	210	212	242	213	2 5	245	218	217	218	219	220	22.4	22.	223	224	225	226	227	228	229	230	231	232	233	234	235	236	237	238	239

Table 2-4: Poor vs. Good Clinical Outcome: Genes upregulated 3 fold or higher in any one tumor sample.

-	-	-	- ,	-	-	-	-	٠ +	- ,		-	<b></b> ·	- •		- •	-		-	-	,	<b>-</b>	- •	- •		- ,	- •	- •	- ,	- ,	- `		-	-	-	-	-	
3.84	1 20		3.35	0.61	0.86	1.70		5 6	17.1	0.89	3.29	<u>1,</u>	1.05	2.80	5.28	3.29	1.52	0.63	3.13	1.14	96.0	0.66	1.47	3.60	0.41	0.63	4.	1.33	0.71	1.30	1.21	4.80	1.39	0.92	3,23	3.61	
1.85		<u>.</u>	1.50	1.46	4.52	40.	0.4	1.38	1.44	3.79	1.70	1.07	<u> </u>	0.16	0.85	1.62	3.89	1.84	2.00	1.22	1:11	1.72	1.19	0.73	0.82	1.26	1.06	1.27	1.17	3.76	3.39	0.90	3.02	1.79	4.1	1.09	
6	3.	1.68	1.70	2.08	8	9.	0.56	1.96	1.38	0.87	1.3	1.23	1.08	4.00	0.73	1.46	0.71	1.62	48.0	1.65	1.46	4:	1.22	1.28	0.58	1.07	0.69	1.27	1.93	1.22	1.27	0.81	98	0.92	0.92	1.26	
,	0.85	3.48	5	• c	5.30	<u>\$</u>	1.20	3.06	3.36	183	3 5	 	3.78	0.76	0.46	0 63 0 63	4.47		2.50	3.33	3.54	3.35	3.26	1.53	5,61	4.15	3.92	3.22	3.27	0.80	1 19	- C	6.50 80 80	0.20	. 6. t	5.5	
	1.89	1 89	5 6	1.80	1.88	1.88	1.88	4 85		1.85	1.85	1.84	<del>1</del> . 3	1.83	1.83	1.83	1.83	1.82	1.82	1.82	1.81	P 6	1.78	1 78	7.0	1.10	178	1,77	177	F		);;;	1.77	1.76	5.7	1.75	1./4
	2.11	: C	D4.	0.89	53.13	2.13	263	20.5	. A	8.26	<b>3</b> 0.	1.38	41.96	0.53	5.23	1.90	1.56	2.99	2.48	2.68	12.19	2.78	2.18	2.31	2.53	0.03	2.80		0.87	80.4 80.4	7.89	2.64	5.15	0.26	7.01	2.78	<b>40</b> .90
	+ +	<u>.</u>	3.97	0.47	28.27	4.14	1 !	<del>.</del> 5	 88.	4.47	0.59	0.75	22.82	0.29	2.86	2.	0.86	<del>2</del> .	1.36	1.48	6.74	1.55	1.22	1.29	1.42	0.01	1.57	4.81	0.49	231	4.46	1.49	2.92	0.14	3.99	1.59	6.25
	7 4 6 7 6 7	161651	AA453531	AA682321	AA443698	640630	AA455565	R98344	AA155787	AA476961	N20989	AA704802	AA684406	AA609982	R38645	AA772497	AA677984	N91744	AA282594	AA434117	AA677397	W88623	AA620421	AA448194	N46759	AA885052	R71124	AA030004	H61082	AA620611	R53342	AA905838	AA827405	AA126682	R70684	AA876354	W05033
		78680	788136	462926	784016	2	813719	201641	590253	739578	265853	452468	868652	1031203	23000	413300	430763	293328	713019	770216	454896	417591	951022	782797	277112	1466485	142944	470216	206172	1048702	40031	1505294	1422794	502593	141815	1493079	298648
																						259										269	270	271	272	273	274

Table 2-4: Poor vs. Good Clinical Outcome: Genes upregulated 3 fold or higher in any one tumor sample.

•		- ,	_	- ,		-	-	-	<del></del> ·				-	-	•	-	-	-	-	<b>-</b>	-	-	-	-	-	-	-	-	<del>-</del> ·	-	-	•	· •	<del>-</del>	-	
Ġ	0.0	36.	1.28	0.48	1.07	3.13	0.81	0.28	3.32	1.21	0.40	0.71	3.71	0.73 E	4.33	3.77	3.29	0.53	3.62	0.84	3.67	0.83	1.05	0.25	0.45	1.25	3.00	1.76	0.16	3.43	1.35	1.17	0.72	66:0	1.24	
;	11.1	0.81	0.93	1.10	1.53	1.38	1.81	0.87	1.21	3.37	1.42	1.42	1.47	1.70	1.21	1.07	1.41	4.89	1.33	1.09	1.42	0.95	68:0	0.76	1.78	1.12	1.46	0.62	0.91	£.	1.31	1.15	1.65	1.23	1.01	
	1.24	1.36	1.28	2.26	0.59	1.60	3.25	2.31	1.20	1.22	1.45	3.33	0.78	1.43	0.78	1.18	0.65	0.51	0.95	1.63	1.03	0.57	1.50	4.85	3.15	1.20	1.11	0.87	2.51	1,01	0.81	1.18	1.21	0.80	1.1	
	3.73	3.43	3.48	3.11	3.75	0.81	1.05	3.45	1.16	1.08	3.62	1.41	0.91	3.01	0.53	0.83	1.50	0.91	0.94	3.27	0.70	4.47	3.29	0.95	1.42	3.17	122	3.52	3.15	96.0	3.23	3.18	3.11	3.66	3.31	
	1.74	1.74	1.74	1.74	1.74	1.73	1.73	1.73	1.72	1.72	1.72	1.72	1.72	1.72	1.71	1.71	1.7.1	1.71	1.71	1.71	1.71	1.70	1.70	1.70	1.70	1.70	1.70	1.69	1.68	1.68	1.68	1.67	1.67	1.67	1.67	O
	2.27	5.31	53	3.02	29.04	2.13	57.00	5.	1.38	0.58	35.33	5.05	1.42	7.12	5.74	69.0	2.60	0.82	1.74	62.95	2.09	0.33	21.50	86.98	3.24	1.78	3.16	90.0	0.07	0.80	10.91	47.59	5.91	8.82	0.85	
	1.30	20.00	2.60	1.74	18.74	12	32.95	3.00	08.0	¥.0	20.62	2.94	0.82	4.15	3.35	0.40	1.52	0.48	10,	36.82	123	0.19	12.62	3.54	181	105	1.86	0.03	9.0	<b>3</b>	6.53	28.46	3.54	5.28	0.51	
	AA417344	T81289	AA457236	AA057032	R09179	AA455080	AA620995	B18943	W71990	N26606	AA459781	AA25551	H80063	WT3724	R81831	AA700843	AA701948		AA464598	AA047260	W86124	R54179	A A 456695	A100302B	H06307	H94563	R82317	H84734	H73808	AA437093	AA464068	AA664389	AA733061	AA420992	AA883114	
	731168	25061	RABBLE	488913	127610	010/31	1056172	2495A	245701	269177	796366	682477	249070	244108	147834	426097	435040	25122	812961	48984	416227	41608	813140	1884774	25580	243159	149058	220135	214713	757325	810284	868630	399562	731255	1466942	
	275	276	277	278 278	270	617	28.4	, ac	283	284	285	286	287	288	280	200	5 5	, 6 6	202	207	204	200	707	200	200	300	5 6	302	303	305	305	306	307	308	308	

3.30 1.22 0.71	1.38	0.67 0.62 3.05	1.68 0.34 0.68	1.58 0.84 5.54	0.94 1.18 0.83	0.85 0.65 0.80 3.88 0.91	0.48 0.70 0.79 0.79 0.79 0.55 1.32	
52.1 80.1 . 05.1	1.06 1.08 3.11	1,45	1.14 1.21 0.75	1.24 1.03 0.28	0.82 1.19 0.58	4.10 1.23 0.94 0.91 3.56	0.88 1.23 1.60 1.04 0.57 0.88 0.41 1.33	
0.92 1.21	0.68 0.61 1.07	3.00	1.57 0.58 2.19	0.59 0.84 0.57	3.28 3.28 1.15	0.55 1.04 1.04 0.86	1.61 1.25 3.05 1.38 1.24 0.81 4.59 0.86	
1.20 3.16 3.86	0.80 3.35 3.30	3.21 3.23 3.23	9.78 3.06 3.22	3.72 9.80 9.80 9.80	3.52 3.52 3.88	3.11 1.00 1.00 1.00 1.00 1.00 1.00	3.39 3.03 3.12 3.26 3.83 0.74 3.04	10
1.67 1.67 1.66	1.66	. 1.66 1.66 1.67 1.67	1.64 1.63 1.63	1.63	1.62 1.62 1.61 1.61	1.61 1.60 1.60 1.59	1.59 1.59 1.58 1.58 1.58 1.57 1.57	
0.23 0.57	2.46 2.82	0.59 4.02 0.80	0.71 0.28 12.12	0.76 9.47 5.76	12.39 17.82 0.89 3.93	3.25 2.10 4.80 0.33 5.92	3.22 2.95 26.88 2.27 7.70 7.70 1.30 1.36 1.75	•
41.0 45.0	25.68 1.48 1.70	0.35	4.83 0.43 7.43	5.83 3.54	7.64 11.02 0.65 2.45	2.02 1.31 3.00 0.21 3.72	2.03 1.86 1.43 1.43 4.88 1.54 7.55 0.86	<u>*</u>
AA884397 7 AA788738	AA983362 AA757417 TE5398 RE9358	H48138 H16439 AA719257	AA479512 AA279990 H15040	H58655 R43553 AA150422	R43053 AA012867 AA469472	AA//4005 W73748 AA487676 AA070435 N48913 AA156926	AA4553500 AA116061 R37357 AA169606 H11036 AA279145 AA456377 H24580	H23548
1 aDI 1461725 1240347	159 <b>2715</b> 395417 21634	193780 47527 1292593	754313 705064 48454	825031 212326 32794	32509 360885 814460	344272 344272 841624 530954 279613 602618	742132 813755 631862 28270 594286 47378 703943 813256 160730	52002
310	312 313 314	316 317 318	319 320 321	322 323 324	326 326 327 328	329 330 331 333	335 335 337 338 338 340 341 343	***************************************

Table 2.4: Poor vs. Good Clinical Outcome. Genes upregulated 3 fold or higher in any one tumor sample.

BNSDOCID 2 MC - 315383640

-	-	-	-	-	-	-	<del>-</del>	-	-	-	<del></del>	-	-	Ψ-	-	-	-	-		-	₩-	-	<b>-</b> -	-	-	-	-	-	-	-	-	-	•	<b>~</b> -	
0.81	3,45	0.47	<b>3</b> .0	3.79	1.26	0.42	0.72	1.26	0.83	0.70	4.14	3.11	3.78	0.85	0.43	3.36	0.22	3.08	0.78	0.80	3.69	3.76	0.31	0.61	0.86	0.69	3.22	97.0	0.56	0.78	0.50	0.77	0.58	0.37	
0.93	1.08	99.0	0.86	0.68	3.21	1.98	0.75	3.08	9.0	1.48	0.56	1.13	0.71	3.38	1.63	0.74	0.76	0.95	0.80	0.67	0.51	0.77	0.52	1.12	0.91	1.14	96.0	0.81	0.59	09:0	0.57	0.71	0.48	0.74	
1.16	0.80	1,45	0.79	0.70	0.71	3.16	8.1	1.08	3.67	3.03	<b>9</b> .54	<del>1</del> .0	0.83	0.85	3.03	1.09	1.24	0.87	0.79	1.03	0.70	0.85	1.55	0.74	3.16	0.76	0.50	67.0	0.87	1.19	3.65	1.03	1.05	1.17	
3.28	0.78	3.59	0.58	1.00	0.99	0.59	3.63	0.71	, 0.64	0.86	0.83	0.78	0.69	0.93	0.78	0.68	3.64	<b>26</b> :0	3.45	3.31	0.86	0.35	3.35	3.26	0.79	3.11	1.02	3.29	3.60	3.04	0.88	3.08	3.47	3.29	
33.	1.56	20.	25.	<u>2</u>	26.	2.	83.1	1.52	1.52	1.52	1.51	1.51	1.50	1.50	1.47	1,47	1.46	1.46	1,45	1.45	4.	1.43	1.43	1.43	1.43	1.43	1.42	14.1	1.40	1.40	1.40	1.40	1,40	1.39	=
2.13	1.22	9.93	3.15	2.24	0.16	1.41	8.37	0.61	2.19	1.42	5.60	2.09	1.58	1.47	60:0	0.73	14.17	1.51	0.16	0.70	1.98	3.43	11.76	2.63	60:0	0.11	4.22	1.23	0.63	16.09	1.11	12.76	0.80	1.83	
1.38	0.79	6.43	705	1.45	0.11	0.92	5.48	9	<u>‡</u>	0.84	3.70	1.39	1.05	0.98	0.0	0.50	9.68	2	0.11	0.48	1.37	2.40	8.21	<u>*</u> .	90:0	0.08	2.96	0.87	0.45	11.47	0.80	9.13	0.57	1.31	
H59188	W24646	AA282159	AA620466	AA400002	T69522	R16541	AA682274	AA779225	R93124	AA886792	AA406320	AA278842	AA912071	N50904	AA161248	H51122	R98472	AA411668	H10073	AA779212	R73088	H02294	H99672	AA644563	AA004671	R01937	H06377	AA677025	R92894	AA193254	W86183	AA428014	N54788	R89082	
204251	308223	712888	951108	743187	66952	129589	461699	452676	196992	1501546	754591	703827	1486752	281045	505414	179631	201125	753271	46844	453988	141548	151184	263697	845521	428796	124567	44300	454564	196612	665774	416305	773483	244310	185751	
345	346	347	348	349	350	351	352	353	354	355	356	357	358	359	360	361	362	363	364	365	366	367	368	369	370	371	372	373	374	375	376	377	378	379	

0.99 3.23 3.78 0.31 0.76 0.70 0.49 0.42 0.42 0.42 0.42 0.42 0.40 0.40 0.26 0.26
0.82 0.52 0.73 3.28 0.49 0.54 0.70 0.59 0.59 0.36 0.36 0.36 0.36 0.36 0.36
3.15 0.80 0.37 1.04 0.78 3.48 3.65 0.70 0.58 0.53 0.53 0.45 0.45 0.45 3.65 0.25 3.40
0.53 0.74 0.78 3.35 0.34 0.66 3.50 0.59 3.08 3.11 0.13 3.10 0.13 3.29 0.12
1.38 1.36 1.36 1.35 1.34 1.32 1.32 1.29 1.29 1.25 1.20 1.09 1.09
3.59 1.84 2.34 14.94 4.30 1.52 3.71 1.41 0.57 1.30 5.27 2.54 7.36 0.10 6.03 2.20 4.23 2.51 14.06
2.61 1.35 1.72 11.02 3.20 1.13 2.79 1.07 0.43 1.01 4.08 7.18 2.03 5.89 0.08 4.98 1.82 3.53 2.3.01 13.33
AA922309 AA706969 W93483 AA698815 T73883 N49574 N49574 N49629 AA102670 AA040265 AA102670 AA0478542 AA478542 AA443706 H49519 AA401137 AA443706 H49519 AA401137 AA465495
1474284 451911 357264 897952 84464 277708 243741 563598 376043 764472 264100 42373 784017 178860 741497 380890 1390584 462325
380 382 382 383 384 386 380 390 391 392 393 398 398 398 398 398 398

Table 2-5; Poor vs. Good Clinical Outcome: Genes upregulated 3 fold or more in at least 2 tumor samples

Tabi4 2-5: Poor vs Good Clinical Outcome: Genes upregulated 3 fold or more in at least 2 tumor samples.
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						7 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 - 1 -	E-14/mm32/	Fold(mpm33/A Fold(mpm34/Av # of Tumor	Fold(mpm34/A)	/ # of Tumor
Order	Clone ID	Accession No.	AveGood	AvePoor	Fold(Averoon AveGood)	AveGood) eGood) AveGood)	AveGood)		(poog)	Samples
,			,	9 <b>)</b> 67	29.71	2.52	57.56	34.98	23.80	က
<b>-</b> ·	753076	AA4304/8	1.42	2	8.08	690	15.64	3.34	3.64	၈
7	124071	R02586	20.0	3.6	9 t	99 6	7.75	1.51	7.73	m
က	346321	W78647	90	3.29			6.82	0.22	6.80	က
4	855624	AA664101	24.18	C6.801	7 7	200	185	4.66	0.57	೯
9	42325	R61073	13.03	27.75	0. t	9 9	8	1.30	3.29	ю
9	287528		2.13	6.32	2.97	00.4	9 6	301	3.60	m
7	756595	AA444051	4. <del>6</del>	3.89	2.67	0.45	19:5	200	57 14	2
ထ	452374	AA700876	0.64	19.98	31.26	0.24	62.30	0.37	1 10	۰ ۱
· თ	357105	W83315	0.87	8.54	9.81	0.81	25.33	12.26	0.85	۷ (
5	460798	AA708152	1.24	6.95	5.61	0.47	17.80	3.83	0.33	<b>4</b> (
= =	284601	N76201	0.27	1.50	5.49	0.82	10.19	0.77	70.77	7 (
: 2	460487	AA677706	53.51	237.74	4,44	0.02	8.87	0.03	8.85 5.05	N 1
i Ĉ	248412	N58558	0.86	3.63	4.25	0.95	7.66	0.75	7.64	7
2 7	212188	H58848	95	2.37	4.23	0.93	6.76	2.47	6.75	7
i t	782543	AAAAAAA	4.08	16.09	3.95	1.06	1.58	6.19	6.98	74
<u>.</u>	782813	AA432030	2 03	7.78	3.83	1.37	1.26	6.00	6.70	2
5 5	129128	R40935	0.38	1.47	3.81	0.92	3.71	0.70	9.82	7
- <del>c</del>	854678	AA630084	9	21.23	3.52	1.40	4.21	2.50	5.95	7
5 5	382059	AA001431	1.67	5.16	3.51	1.66	5.87	99.0	5.86	2
2 5	454822	AA677388	0.60	2.09	3.47	0.37	9.64	0.26	6.62	7
3 2	308696	N92842	0.05	0.16	3.28	1.59	6.55	0.00	4.97	8
3 5	416766	W84751	1.20	3.83	3.19	0.75	5.82	0.36	5.81	7
1 8	489535	AA099252	3.91	12.41	3.17	5.95	0.84	1.37	4.54	7
3 2	264.865	AA028171	08.0	2.53	3.17	3.39	2.18	4.93	2.18	7
, ,	740047	AA478298	53.79	169.35	3.15	1.55	5.32	0.41	5.31	7
3 6	241705	H01580	171	5.18	3.04	1.25	4.71	5.51	0.69	8
7 2	754559	AA406301	25	3.96	3.00	2.54	3.88	4.35	1.25	2
, c	487497	AA045524	0.49	1.46	2.96	1.51	4.72	0.91	4.72	8
0,00	843098	AA488676	5.29	5.55	2.94	1.27	2.66	3.34	4.49	8
3 E	298303	8582N	0.22	0.65	2.94	0.99	4.79	1.18	4.79	7
3				:	•					

папапапапапапапапапапапапа
4.42 4.42 4.95 0.00 3.66 3.81 2.47 3.92 3.19 3.18 3.18 3.18 3.19 3.19 3.19 3.19 3.19 3.19 3.19
4.07 4.85 0.29 0.83 2.12 2.04 1.06 1.06 1.06 1.06 0.94 3.21 3.21 3.28 3.21 3.09 0.53 0.53 0.34 0.53 0.53 0.34 0.53 0.35 0.36 0.36
1.80 0.86 5.28 4.59 3.14 3.67 3.82 1.22 1.22 1.25 3.22 3.23 3.34 3.39 3.39 3.39 3.49 3.39 3.49 3.49 3.49 3.49 3.49 3.49 3.49 3.49 3.49 3.49 3.49 3.49 3.49 3.49 3.49 3.49 3.49 3.49 3.49 3.49 3.49 3.49 3.49 3.49 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55 3.55
1,32 0,83 0,62 1,13 1,19 1,19 1,13 1,14 1,11 1,13 1,13 1,13 1,13 1,14 0,57 0,39 0,39 0,42 0,42 0,39 0,42 0,42 0,57
2.90 2.87 2.86 2.67 2.64 2.56 2.56 2.34 2.34 2.34 2.34 2.34 2.34 2.34 2.34
4,08 4,28 3,16 1,86 0,02 43,42 1,70 3,00 19,45 10,19 60,88 69,88 4,88 4,88 4,15 0,13 1,05 1,05 1,05 1,05 1,05 1,05 1,05 1,05
1.41 1.48 1.10 0.67 0.01 1.20 1.17 7.79 6.54 0.08 3.43 3.39 29.86 0.55 2.09 20.98 1.81 0.06 0.49 0.21 2.05 0.49 0.21 0.06
AA432049 WA6900 AA682386 AA436317 AA436317 AA436317 AA48568 AA701519 AA611215 AA011215 AA011215 AA011215 AA014499 WA2812 AA041499 WA2812 AA041499 WA2812 AA4350278 R53558 R53558 R63546 H1063 AA452278 AA461174 H86564 AA001441
815542 782537 324437 461759 754393 770883 415892 435820 823696 811000 754600 41214 898096 1473304 323238 770880 303099 39959 138788 52071 47418 787938 787938
25 25 25 25 25 25 25 25 25 25 25 25 25 2

Table 2-6: Poor vs. Good Clinical Outcome: Genes upregulated 5 fold or more in at least 2 tumor samples.

Table 2-6: Poor vs Good Clinical Outcome: Genes upregulated 5 fold or more in at least 2 tumor samples

. ŭ	IMAGE Clone Id	Accession No.	Tissue Distribution	AveGood	AvePoor	Fold(AvePoortA Fold(mpm31/Av Fold(mpm32/ Fold(mpm33/A veGood) veGood) veGood)	Fold(mpm31/Av •Good)	Fold(mpm32/ AveGood)	Fold(mpm33/A veGood)	Fold(mpm34/ AveGood)	# of Tu
75	753076	AA438479	thymus	1.42	47 CF		4				
45	77774	3700076			2	7.87	7.52	57.56	34.98	23.80	
? ;	100	0/900/20	prostate	20.	19.98	31.26	0.24	62.30	72.0	7, 0	
35	27105	W93315		0.87	8.54	186	, a	20.00	70.0	02.14	
*	4601	N76201	prain	76.0	1		5 6	3	12.26	98.0	
Ř	6321	W74079	000	1 0	00.1	9.48	0.62	10.19	0.77	10.17	_
, K	£624	0.0000		4	3.29	5.17	3.69	7.75	1.51	7.73	•
3		101 #00 <b>%</b>	lestis	24.18	108.55	4.49	4.12	6.82	66.0		. ,
<b>\$</b>	0487	AA677706	poue	53.51	237.74	4 44	.00		7	0.80	•
246	<b>B4</b> 12	N58558	liver	0.88	163		0.02	0.0	0.03	8.85	•
212	2188	H69148	liver	98.0	9 6	C7 .	C. 8.7	7.66	0.75	7.64	
782	2513	AA432030	1	D 6	75.7	4.23	0.93	6.76	2.47	8.75	"
7.87	1813	A 4 423030	\$11\$91	3.	16.09	3.95	1.06	1.58	6.19	89.8	
9 6		AA432030	lestis	2.03	7.78	3.83	1.37	1.26	00		• (
205		AA001431	pone	1.47	5.16	3.51	9	. 20 3		0.70	~
454	(822	AA677388	colon	0.60	200	3.47	3 6	5.07	99.0	5.86	~
415	766	W84751	heart	1 20	3 6	÷ ;	0.3/	6.64	0.26	6.62	7
740	740941	AA478298	factle	27 23	3 5	91.0 -: 0	0.75	5.82	0.36	5.81	7
30.0	437	WARDON		20.50	109.33	3.15	1.55	5.32	0.41	5.31	•
•	ì	0000	Done	1.10	3.16	2.86	0.62	200	•	•	•

Table 2-7: Good vs. Poor Clinical Outcome: Genes upregulated 3 fold or more in at least 1 tumor sample.

Table 2-7: Good vs Poor Clinical Outcome: Genes upregulated 3 fold or more in at least 1

Samples	4	4	٣	en	. «	<b>,</b>	, es	n	က	က	က	က	e	•	~	) «	י נ	<b>n</b> (	- O-	m ·	ന	n	က	က	က	<b>6</b> 0	7	7	7	7	8	7	7	7	7	7	7	7	. 6	1	
old(mpm36/AV ePoor)	7.30	6.26	24.45	2 0	00.	8.80 08.30	15.90	10.17	5.86	12 92	3.28	5.47	7. 6	- o	9 7	E.9.	5.03	6.23	0.54	1.35	3.77	3.74	3.11	3.97	3.28	3.31	1.02	1.22	3.17	0.71	7.09	2.65	7.23	10.1	0.98	0.31	3.00	0 0	b 5	r n	
old(mpm30/Av F aPoor)		0.68	<u>-</u> :	2.10	9.83	7.05	0.52		0.00	9 9	, 4. c	30.00 00.00	3.86	2.14	<b>.</b> 8	4.30	2.55	3.51	2.28	1.32	127	476	25.	2.0	332	3.13	4.49	1.65	5 2	5 5	58.0	7	 	200	68.0		7	9	0.93	2.00	
Fold(AveGood/A Fold(mpm27/Av Fold(mpm28/Av Fold(mpm30/Av F		3.51	6.33	4.22	0.27	73.0	20.0	05.1	0.98	9	0.35	0.80	0.46	4.46	0.79	£.	75	, a	0.00	9. C	77.7	1.42	<b>\$</b> {	2 5	3.67	કું કુ	8 S	26.4	4. 5 5. 5	33.56	15.74	CR.7	5.5	5.0 5.0 5.0 5.0 5.0 5.0 5.0 5.0 5.0 5.0	10.09	11.86	11.75	10.11	10.03	0.68	
old(mpm28/Av l		6.80	3.61	000	24.	<del>.</del>	0.41	1.97	1.98	290	3	130	0.35	253	F: 02	<u> </u>	~ (• • :	2.38	1.45	4.18	4.22	0.87	1.71	3.69	3.21	1.66	1.20	25.69	7.47	1.15	5.93	2.75	3.53	2.3	4.59	4.36	5.86	3.36	5.41	0.55	}
Fold(mpm27/Av F	eroor)	707		2.5	21.13	22.87	22.41	8.50	14.00	12.05	6.03	5 3	17.10	80.41 10.41	)O.8	12.66	3.28	4.86	6.03	5.36	5.71	7.31	5.86	2.50	2.22	4.16	3.23	99.	1,65	4.	1.46	9.05	2.55	2.64	0.50	00	0.36	89.0	9 G	2 5	3.62 4
Fold(AveGood/A	vePoor)	,	2.0.	4.21	10.17	8 29	7 87	0.7	4.	5.48	5.78	5.50	5.03	€.90	4.38	4.16	3.83	3.87	262	3.43	3.5	3.30	9.5.6 8.0.6	200	1 26	276	75.0	11.65	4 29	4: «	8 5	9 ¥	S. 4	9 6	- T	5 6	0.6	37.6	3.68	3.60	3.65
Aughor	Averog		0.50	6.85	8	3 5	7	3.45	0.45	0.42	4.62	6.9 18.0	0.82	0.71	500	5	5	8.5	9 6	0.0	3 5	25.99	8	77.	<b>3</b> 8	27.00	<b>*</b> !	2.5	0.37	9.0	D8:0	0.40	3 6	8; :	÷ !	0.67	0.32	0.75	0.53	9.0	0.49
	AveGood		2.55	78.87		20.	38.67	27.13	3.32	2.70	26.72	38.22	4.13	07.5		3 8	3 5	6.27	2.50	<u>+</u>	<b>-1</b> 84.	87.41	5.16	4.16	0.12	76.12	702	17.01	4.28	10.81	7.27	1.41	<b>4</b> .0	32.22	5.53	2.58	1.22	2.84	19.	3.43	1.80
Gen Bank	Accession	Number	WOS628	A A 486471		2505	N47117	N48259	R71393	H65286	AA430545	AA282273	TR0111	NEGIES		20000	LACK .	H43317	AA778675	R88904	H16743	<b>AA456975</b>	N45282	AA626248	H15926	AA779728	R68272	AA777187	AA424535	N95226	AA134576	R66438	N90882	R35051	AA001219	AA443903	N21084	AA668929	AA025434	AA424920	W94121
	Clone			A JOART	811162	471641	281039	282089	465072	2000	2007	7,000	DCR7U	1551	28//28	41228	122345	186918	1049033	195365	50114	838611	283196	745525	159487	1034473	138444	378488	787130	307337	502625	41869	301735	136805	362278	756708	265028	4400509	765707	769063	147628
	Order	<b>i</b>		-	7	n	4	r u	n a	0 1	~ •	xo (	<b>о</b> ъ '	5	Ξ	12	13	<b>*</b>	45	4	: =	: #	2	2	7	2	12	7	25	8	27	28	78 78	8	8 2	÷	; ;	3 3	\$ ;	2	3 :

Table 2-7: Good vs. Poor Clinical Outcome: Genes upregulated 3 fold or more in at least 1 tumor sample.

2	7	7	2		• •		• •	• •	<b>4</b> (	<b>v</b>	7	7	7	74	7	7	7	7	74	7	2	. ~		4 6	٠,	<b>v</b> (	۰,	7	7	71	7	7 (	7 (	7 (	7 (	4 6	٠,	4 6	4 (	v (	<b>4</b> (	<b>,</b>	7	~	7	7		
1.38	77.6	8 17	4 57	, c	, d	9 6	1 0	90.0	0.35	\$ 5°	3.69	3.25	<b>9</b> .18	1.08	6.69	0.75	9.12	1.75	4.74	5.77	2 19	4.74	<u> </u>	* 6	70.	4.6	<b>4</b>	0.72	1.24	0.76	<b>4.8</b> 9	2.27	3.42	6.33	5 8 8	 	- 6	8.4 8.5	0.0	3 8	3.28	8.	\$	1.12	3.25	6.89		
<u>4</u>	5.51	70.6	6.0	0.90	4/.7	מ מ מ	C:7	09.1	90.0	2.50	2.35	2.42	1.32	0.66	0.49	3.20	09:0	1.70	6.81	4.53		8.6	20.0	0.00 1.00	1.17	6.48	2.55	0.46	2.85	69.0	1.56	3.01	2.18	.00 .00	<b>5</b>	3.15	3.69	5.65	1.21	1.21	2.49	1.76	6.54	1.13	1.58	0.28		
2.07	5,5	2.73	25.1	2.38	2009	10.28	0.51	1.61	2.26	1.26	0.59	0.91	0.45	1.80	1.77	0.57	0.74	4.46	0 04		20 20	3.16 0.50	27.9	1.02	0.79	1.14	1.55	7.31	4.22	7.85	1.62	1.71	9.54	0.65	6.17	5.37	2.58	8.	2.67	5.03	2.12	1.95	0.61	5.61	1.56	0.22		
7.0		0.52	<b>5</b> 7.7	16.	2.54	3.45	1.09	4.91	3.72	2.96	1.26	1.48	0.81	5.84	0.71	0.89	4 10	2 99		2 .	4.	6.21	3.75	3.46	1.98	0.96	1.25	4.28	2.02	5.0	1.22	2.68	1.46	0.37	1.13	0.86	1.20	0.83	3.19	4.91	3.72	3.10	0.53	3.38	2.03	3.78		
0	<u>n</u> ;	1.4/	.63	1.01	3.72	0.86	8.38	0.59	8.99	6.07	8.02	7.62	3.86	6.21	368	10.04	0.74	2.50	3	0.40 0.40	0.82	0.92	1.83	6.94	4.81	0.58	4.87	0.63	3.01	3.32	3.79	3.41	5.42	1.63	3.54	2.51	2.17	0.44	1.63	0.43	0.61	3.24	0.31	0.80	3.47	0.70	2	•
č	\$	3.53	3.51	3.36	3.35	3.29	3.29	3.28	3.26	3.23	3.18	44.5	3.12	712		. 6	6 6	8 8	7.00	2.81	2.81	2.81	2.77	2.75	2.73	2.72	2.71	268	2.67	2.63	2.62	2.62	2.60	2.60	2.59	2.59	2.55	2.50	2.46	2.44	2.44	2.41	2.41	2.41	2.38	2.37		
;	7.90	8.55	0.02	<b>4</b> .	1.14	0.79	3.68	2.46	0.25	1 18	,,,	77.7	86.0		3 6	20.0	9 6	7.5d	3	36.00	0.75	35.15	0.43	0.83	17.94	14.62	122	g	261	75.0	8	50	1.17	2.46	3,	3.	0.32	15.73	7.91	0.77	138	0.48	21 78	0.67	1.52	8	! !	
1	28.75	30.23	90.0	13.79	3.83	2.60	12.10	8.05	0.83	2 7 9		60.7	90.6	8 5	5 6	9 6	7.75	4.17	7.87	101.29	2.12	98.80	1.23	2.29	48 94	39.72	331	60.0	90	, t	282	8	20.00	90,00	244	4.21	0.81	39.31	19.49	1.87	3.37	1.16	52.42	1 37	3.61	3.93	<u>;</u>	
!	R45192	R77948	AA443967	AA633658	AA412417	R68101	H09076	R00276	R61372	AA678113	N.53447	12C3T7.0.0	TE7128	04432270	04707E	0 / 0 / 0 / 0 / 0 / 0 / 0 / 0 / 0 / 0 /	705057	AA416/40	K4551	N55459	N59816	AA196979	AA700680	H92821	AA487902	T56281	AA878120	T69477	AA496988	N53011	AA150500	T69305	W74533	N74131	H17121	AA702335	R61883	H72722	N34316	AA453526	AA634431	R54590	H53340	N23139	AA479795	H73590	•	
	35058	145740	757200	856575	731469	140197	46168	127264	17080	4055707	VD/5501	117647	0800/9	ACCOS.	CCC70/	190081	1/4101	768020	35328	245990	288983	646037	434952	271574	840590	78353	1418621	58902	823658	00000	401754	82905	746583	298417	51178	447418	42872	232772	277173	795425	743868	40100	202636	267548	740604	214441	: :	
	38	39	\$	4	: 2	<b>4</b>	: 4	. 4	<b>?</b>	} ;	÷ ;	<b>5</b> 6	P	3 3	ភូមិ	8 8	3	<b>3</b> :	S	જ	22	58	59	8	3 2	; £	, E	3 2	<b>5</b> 4	3 4	3 2	S &	3 2	3 2	: =	2	R	7	22	292	1		9 9		} <b>=</b>	- 2	<b>,</b>	

Table 2-7: Good vs. Poor Clinical Outcome: Genes upregulated 3 fold or more in at least 1 tumor sample.

ŗ	4 (	7 (	7	7	2	•	۰,	<b>4</b> (	<b>4</b> (	7	7	7	7	7	7	0	۰ ۱	, ,	4 6	4 (	4 (	7 (	7	7	7	74	8	8	7	7	8	7	2	7	2	7	٠ ٢٦	7	7	7	7	7	7	7	ď	8	~			
!	3.86	1.24	4.33	4 96		۲ ر د	 	0.90	1.87	4.40	3.90	1.32	5.80	20.4	50.	90.4	<b>9</b> . 4	3.83	1.53	3.95	0.92	3.62	3.44	4.38	3.80	1.03	3.61	75	3 O.E.		80	330	3.07	3.11	3,16	3,85	4.12	4.24	3.47	3.05	0.61	3.76	3.15	<u>=</u> :	0.61	0.49	0.48			
	2.42	3.44		26.7	3.80	0.18	4.10	0.74	69.4	0.24	02.4		3 3	16.0 1	5.78	5.39	0.23	<del>1</del> .06	0.71	5.39	1.60	1.67	203	3.46	) t	50.	3 3	4 2	\$ 6 \$ 6	2.36	0.40	. o.	9.19	200	0.95	, ,	25.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00 15.00	. 65.0	) (F	2 5	27.50	2 6	3.37	2 40		6.0	4 50	} ;		
	37.0	2 5	7.47	0.53	1.05	0.08	5	3.5		24.0	. o	0.91	2.15	0.45	0.43	0.59	0.25	-	20.0	. OF C	, c	) (4)	3.00	0.93	1.22	79.0	3.02	1.80	0.49	0.41	1.13	0.52	0.98	1.23	0.68	0.93	0.78	0.57	0.38	0.65	1.87	c0.4	0.52	0.21	3.72	3.89	3.81	3.25		
		1.42	1.57	0.44	4 4	2 6	7.10	0.65	64.	0.83	5.59	10.	0.82	0.78	 		70.0	4.4	6.0	3.49	0.49	3.05	1.35	0.50	0.67	0.93	1.92	38	2.55	3.29	3.70	95.0	69:	1.05	1.99	0.93	0.75	3.84	3.02	0.90	3.19	3.33	99'0	<del>_</del>	99.0	3,19	3.18	0.53		
		3.36	70 %	, c	3.36	3.12	4.65	2.81	4.73	3.34	0.76	9 9	5, 6	97.0	3.40	0.17	0.32	1.03	3.28	4.01	0.34	1.59	0.89	3.64	0.76	90.0	3.33	0.50 Fa C	, c	90.7	90°C	5.50	0.75	2.5	3.13	1 39	1.40	0.92	1.25	0.74	0.91	0.55	0.42	0.69	0.26	0.57	0.36	0.00	က	,
		35 6	3	2.32	2.31	2.30	2.27	2.26	,	3 6	3 6	77.7	2.22	2.20	2.19	2.19	2.18	2.17	2.17	. 4			7 7	1.7	01.2	2.10	2.09	2.07	2.06	5.06	2.04	2.01	66.	<del>1</del> .99	1.98	16.1	1.90	90.4	De: 4	9.0	50°F	0.0.1	0. F		 •	7.7.	9/:	 	20.	
		•	 55.	0.33	40	77.0	5 6	6.0	07.1	0.70	2.7	12.67	75.52	1.40	17 15	15.57	96.6	3 6	08.7	77.0	). (2.6)	10.30 30.30	1.71	3.65	22.41	0.86	0.30	0.83	10.94	9.27	3.81	0.0	25. <b>4</b>	2.07	1.1	1.50	0.85	1.51	2.19	8.31	0.76	15.04	13.84	3.24	21.21	27.34	324	9. 9. 9.	70.0	
			3.15	7.0		7	\$ 5	15.54	2.85	1.57	4.78	28.11	12.27		2.5	20.10	3.5	29.10	6.47	0.52	3.35	22.38	3.60	7.69	47.09	1.81	29.0	1.73	22.57	19 09	7.78	0.02	50.71	4.12	2.20	2.96	1.67	2.97	4.29	15.71	4.	28.35	25.74	5.85	37.58	48.33	5.72	1.59	0.03	
1 200			AA021586	197030	60000	84441524	H08852	T67053	N52151	AA456299	AA456321	AA663981	A A G 8 B 3 4 K	20002	K/624/	H15803	H777765	N57872	AA884926	H50086	AA434487	R16596	W48563	N72137	AA070226	AA779467	H15877	A A 609568	AA111969	A A REBROT	A A R 3520	A 4 4 5 9 2 9 2	N80129	AA035558	R36889	AA459197	AA040424	N64706	N92646	770057	AA457528	W70234	N30680	AA489666	AA630006	R27004	AA418000	R46000	AA609914	
			000100	30405	295985	782575	45852	66560	284383	242154	943470	013178	800/40	1605539	144747	50772	214162	247117	1468260	179083	770868	129585	324R15	204007	ZB1037	230014	1032403	49303	וט/ רגטו	530185	853998	378813	810899	785/87	4/1082 37008	040873	485854	293292	288337	80048	838500	743987	258300	824340	884606	142702	767449	35804	1031113	
				3	\$	85	88	8 &	; a	8 8	20 8	3	5	23	8	3	50	8	6	. S	8	\$ 5	3 3	5 9	2 3	<u>s</u>	\$ 1	50	\$	<b>1</b> 07	<u>\$</u>	<u>6</u>	2	= :	<b>1</b> 25	2 ;	* ;	- 4	- +	- 9	2 2	= 5	<u> </u>		<u> </u>	3 5	125	<u> </u>	127	

<del>-</del> -	-	-	_	<b>,</b>	-	-	-	-	-	-	•-	-	-	-	-	-	-	-	-	-	<b>-</b>	-	-	-	-	-	-	<b>-</b>	-		<b>-</b> - ,	- ,		- •		- ,				-	-				
1.08 0.98	0.54	0.91	0.47	0.47	14.57	15.05	0.83	1.02	0.28	0.95	1.56	06.0	2.81	1.20	<del>*</del> :	0.25	1.53	8.39	1,63	2.96	53	9.72	0.46	1.25	1.22	1.08	0.64	2.10	1.97	1.16	1.15	<del>.</del> 8	0.78	76.4 10.1	7.29	68.0	1.24	1.09	2.20	1.37	8.56	0.99	2.52	i	
1.75	0.77	2.45	1.00	0.72	96.0	1.52	0.98	0.81	0.30	8	11.98	0.82	1.78	0.92	0.97	0.14	0.45	4.	, c	2.51	2.2	207	0.32	980	1.74	53	\$	1.63	2.72	1.19	0.47	1.01	1.31	2.94	2.45	2.34	1.38	6.91	0.84	0.88	151	0.80	1.08	•	
112.76	29.22	0.71	22.44	20.07	2.10	66.0	15.25	1 03	710	2 7	570	2.0		50 GZ	1 28	12.03	3 0	) <del>.</del>	<u>.</u>	47.1	9 C	6.33 0.63	08.0	0.03	7.54	70.8	9.78	1.37	4.50	2.45	1.38	1.49	1.46	1.29	1.37	6.98	1.81	2.31	0.72	1.86	0.82	108	50.7	5.	
0.74	72.0	5.5	5 6	5.5	5.		65.7 87.7	÷ ;	7.75	77.0	75.1	14.7	2.30 2.78	60.0	20.0	47.7 7.0	0 5	1.53	1.41	2.08	1.78	1.67	9.0	. c. c	4.24	 	<u> </u>	1.65	1.92	1.71	2.48	1.62	2.56	2.09	1.06	0.82	0.60	1.66	0.45	184	5 5	5.6	7.00	/c.r	
0.70	 	1.41	75.57	0.43	26.0	1.25	0 6	1.32	19.18	18.72	1.57	2.12	10.78	0.50	07.7	9.47	76.0	11.19	1.61	7.23	6.18	2.06	0.85	11.85	9.16	2.20	~ ·	- F	55.5	1 .	. 80	5.37	7.33	2 18	1.23	2.26	8.17	, <del>,</del>	7.00	2 3	7.04	0.80	7.88	6.15	4
23.41	6.60	6.50	9.90	5.09	90.4 90.4	40.4	3.99	3.97	3.95	3. 26.	3.89	3.71	3.59	3.48	3.25	3.08	3.03	3.00	2.97	2.90	2.86	2.86	2.84	2.83	2.82	2.81	2.79	2.77	7.7	5.75	27.7	2.70	2.59	268	2.68	99.0	2,00	5 6	5.65		2.60	2.57	2.55	2.53	
0.85	0.69	1.23	0.50	0.90	<u> </u>	1,11	0.38	0.36	0.49	3.35	0.50	2.97	5.86	4.72	0.47	0.83	5.90	3.77	2.50	0.62	1.86	0.40	14.1	1.98	66.0	1.58	0.55	1.73	0.87	26.U	24.0	2 6	8 CC	6 6	0.03	0 50	10.20	2 6		<b>5</b> ;	0.58	0.57	0.58	60:0	
19.97	4.58	8.01	2.77	4.56	6.75	4.47	1.51	1.43	1.93	13.18	1.95	11.01	21.03	16.43	<u>7</u> .	2.58	17.91	11.29	7.42	1.79	, sc.	1.1	4.00	5.61	2.78	4.39	<u></u>	4.81	2.40	.43	<del>-</del> ;	<u>-</u> :	e	9 6	9 6	807	27.30 20.00	7.3	1.35	34.0	1.51	1.48	1.43	0.24	
N78927	N48325	AA702640	90806N	AA677574	AA670123	AA487385	AA001834	R93176	N30328	AA845158	AA464880	AA402874	AA454018	R28397	AA182848	AA456008	AA680136	AA680186	A1005521	R66006	H52001	W81198	R00284	AA452113	AA398348	H29897	H20046	AA018338	H45810	H11433	AA504842	A204854	M400183	20000	N9440/	*S0C100	2029AA	CACADA 3	AA020/3/	A1 /90%	H80853	AA680070	<b>T99617</b>	R44664	
300051	279690	384015	303139	455271	844703	841507	427978	275738	258063	1412481	838856	741831	795265	133130	611581	812105	433155	430465	1637302	140131	107525	347429	123265	786525	726835	190325	172785	362419	188151	47652	825736	825719	753770	757907	309893	360213	416374	228462	1049287	293078	230317	430368	123354	34129	
_	_	_	31	132	133	ठ	135	98	(37	88	65	3	<del>-</del>	142	143	<u> </u>	571	148	2 2		2 0	50	55	52	153	7	155		157	_	28	_	<b>.</b>	70	63	3	65	8	22	891	691	20	17	72	

-	-	-	-	. 🕶	- •	- +		<b>,</b> -	Ψ-	<b>-</b> -	-	-	•-	-	. 🕶		. 🕶		. 🕶					•	· <del>-</del>	-	· <b></b>	_	<del>-</del>	-	<b>,-</b>	<del>-</del>	<b>-</b>	-	<b>-</b>	-	-	-	<b>-</b> ·	_	,	- •	- •	- •	-		
1 94	76.0	78 0	5 6	8.5.5 0.0	0.36	1.95	1.55	3.31	2.62	0.58	90.6	4.02	1 0	6 C	76.1	1.4.	ō	5.	55.	2.16	1,18	1.63	0.91	57.1	1.7.7	7C.1	0.83	, ,		67.1	1.30	2.01	 	6.0	2.83	181	3.70	0.10	1.01	0.84	1.73	1.12	1.62	1.26	1.32		
;	0.60	25.7	18.0	3.69	0.87	1.37	1 83	7.4		0.13	707	3.29	2.65	0.58	0.68	0.50	<u>4</u> .	1.02	2.50	1.77	2.96	1.45	0.73	1.20	65.0	2.58	0.86	1.7.1	2.47	0.98	2.21	2.15	2.24	<del>2</del> .	2.27	2.46	1.80	 	0.70	3	3.43	40.	2.30	0.75	2.72		
	6.37	1.21	5.79	60		5.5	, i	5.6	2.59	1,31	2.07	2.31	5.23	2.79	1 29	171	3.50	80	2 14	4 47	2 07	5	2.5	8.8	09'0	4.41	8.86	0.38	3.50	5.01	0.92	2.37	1.22	0.59	1,38	1.73	1.26	2.07	1.50	CL.1	47.7	7.50	8. e	5 5	28	3	
	1.73		, i	2.70	 	8.09	0.87	5.1	2.05	2.57	79	, .	2 9	1.43	77	0.74	2.44	3.0	2.83	18.0	89. 1 100. 1	1.57	28.	9 5	2.73	÷ 6	7.65	S 4	8 5	8 6	8.7 F	5	9.3	78.0	2	0.92	2.38	2.45	4.49	96:0	<u>.</u>	55	1.78	2.64	2.85	8	
	9	2.48	5.32	2.25	2.94	1.39	6.85	1.73	09.0	6.7 7	- 0	0.42	2.81	1.48	6.94	2.00	6.15	2.76	6.14	5.00	2.98	3.34	1.30	8.49	1.00	6.20	1.83	0.40	2.79	1.31	1.87	6.37	1.24	3.84	7.58	J.80	3.02	4.2	2.81	7.79	7	2.87	6.09	1.86	5.46	3.85	ഹ
		2.50	2.50	2.49	97 6	04.7	0.40	2.48	2.48	2.47	2.47	2.47	2.46	2.45	2.45	2.45	44	2.44	2.43	2.42	2.41	2.40	2.40	2.40	2.40	2.39	2,39	2.38	2.38	2.38	2.38	2.37	2.37	2.36	2.36	2.36	2.35	2.35	2.34	7.32	7.37	2.3	231	2.30	2.29	2.29	
		9 0	2 6	9.0	7.40	1.86	96.0	2.05	0.25	1.02	0.02	1 70	10.4	77.1	, .	5.5	3.12	0.03	2.13	0.82	0.9	0.90	2.5	\$ F	£	9 0	. e	2	- 6	2 6	. 4 A B	- F	2	12.22	0.42	2.54	0.67	0.62	<b>0</b> .08	<b>6</b> .0	0.29	0.59	2.31	8.5	9.14	. c.	•
		į	4.5	1.99	6.18	4.61	2.42	5.08	63	3 6	3 6	5 6	4.40	3.11	3.62	24.84	7.63	0.12	5.18	1.98	0.74	2.17	2.93	3.23	6.57	1.40	2.08	35.	<b>‡</b> 5	G 6	20 C	<b>7</b> €	2 8	0.U0	6.63	- K	1.57	1.46	0.19	0.10	0.68	1.37	5.33	3.83	18.76	0.79	ò o
7 2000			AA455654	AA047340	AA437099	AA953357	AA609218	H09859	A 406681	AA486301	AA207124	<b>AA</b> 262080	AA875813	AA464143	R03980	R39891	N57002	AA620783	AA455087	AA444083	AA039851	AA707085	AA431887	AA035450	H79534	R49731	N72115	T64312	AA029578	AA13661B	R39325	AA190789	AA425947	AA666348	AA045595	H03003	AA8/2332 H15099	N50853	W74653	N33264	R38891	AA872006	AA405543	R27193	H69691	N91385	W86653
			814001	509516	75727	101001	1070701	10314/10	4636/	840967	701819	685801	1492202	810328	128738	26736	2777596	1049321	812294	756657	375827	451587	773639	471725	239611	38554	291057	83920	368859	490946	24176	627351	760299	858849	509731	137984	1584628	40004	280957	272501	25052	1475648	772418	133341	212784	306013	416833
			173		. !	e i	178	111	178	179	180	181	182	81	<u> </u>	484	48	487	. e	2 4	9	2 7	- CO	10.	<u> </u>	195	196	187	198	199	8	ຂ	202	203	204	205	508	8	88	508	220	17 6	4 5	3.5	2.5	218	217

Table 2-7: Good vs. Poor Clinical Outcome: Genes upregulated 3 fold or more in at least 1 tumor sample.

1.46 6.91 1.85 0.64 1.21 1.73	2.1.18 1.18 1.13 1.03 1.03 1.03 1.03 1.03 1.03 1.03	2.44 2.44 1.20 1.31 1.32 0.91 2.06	2.13 0.78 0.70 0.70 1.92 0.63 3.15 2.49	2.24 3.11 2.08 2.31 0.65 1.14
2.16 1.63 1.96 0.63 1.03 1.83	1.84 1.23 1.52 1.14 2.03 2.03 1.44 1.98 1.67	1.52 2.03 1.89 1.66 1.66 3.94	2.31 1.54 2.20 3.68 1.68 0.75 0.75	2.02 1.32 1.07 1.75 0.93 1.74 0.63
0.63 1.40 1.50 1.50 1.50 1.50 1.50	5.54 5.46 5.42 5.42 1.20 1.20 2.73 2.11	4.87 0.79 2.87 4.67 1.83 0.94 6.28 5.05	1.18 5.62 1.78 0.93 1.05 1.05 1.03 1.54	2.76 1.64 2.05 1.60 3.22 3.22 0.84 1.49
6.23 1.34 2.73 2.76 1.95 1.95	1.85 0.71 0.71 2.44 1.67 2.58 1.59 0.89	2.13 2.13 2.13 1.46 1.27 2.37 0.60	2.63 1.64 1.64 2.63 1.61 1.61 1.78	1.12 2.37 1.39 1.33 1.57 0.71
1.11 0.84 0.61 6.34 5.62 2.17 2.63	0.70 7.25 1.54 1.18 1.13 1.45 1.06 0.96 3.76	2.71 2.71 3.58 0.90 5.62 0.82 0.67 1.76	2.13 1.05 1.05 2.03 2.48 0.86 1.70 1.71 2.42 1.25	1.54 3.08 2.25 1.42 5.74 1.00 7.07 5.29
2.27 2.26 2.25 2.25 2.25 2.25	2 2 2 2 2 2 2 3 3 4 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5	7	7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7	2.13 2.13 2.13 2.13 2.12 2.12 1.12
0.02 0.90 1.76 0.92 0.84 0.15	0.42 0.42 0.42 2.64 1.13 0.74	9.00 9.00 9.42 9.42 9.43 9.43 9.43 9.43	0.21 0.06 0.06 0.54 0.54 1.57 1.57 1.67 4.96	0.57 29.69 0.07 0.97 1.63 2.88 0.34 1.14
2.05 2.05 3.97 2.07 1.89 0.35	13.95 1.56 0.95 0.44 2.7.25 3.86 1.62 1.62	4.00 6.00 6.00 6.00 6.00 6.00 6.00 6.00	0.48 6.27 0.13 0.13 1.16 1.16 1.35 1.35 10.58	5.21 63.20 0.14 2.07 3.46 6.11 0.72 2.41
AA010619 N29914 AA044906 H40921 AA703652 N86572	H54417 AA460422 AA876265 H10342 T72067 AA703396 R00884 AA148862 AA054135	AA663092 H08397 AA647560 AA055221 AA005350 H10036 W02426 AA460313 AA432253	K38179 NS3167 AA57138 H19324 H72093 AA608575 AA137078 H02039 AA26275 AA708438	H15408 AA156054 H12081 AA706738 AA67468 AA67469 AA074049 AA056013 T60063
430336 271050 488706 177074 450382 278808 814662	203003 796480 431553 46958 85800 450049 123971 566440 380437	452677 45501 841664 377368 428377 48820 296170 785755	23869 245547 810458 60922 213604 960710 491196 151087 745072 506011	49227 590338 47597 1239995 641621 864662 363189 377692 81320
218 220 221 222 223 224		234 235 237 238 240 241	25 25 25 25 25 25 25 25 25 25 25 25 25 2	25 25 25 25 25 25 25 25 25 25 25 25 25 2

Table 2-7; Good vs. Poor Clinical Outcome: Genes upregulated 3 fold or more in at least 1 tumor sample.

-	-	-	<b></b> .	-	-	-	-	-	-	-	-	-	-	<b>,</b>	-	-	•			- •	- ,	- •	-	-	-	-	-	-	-	-	-	-	, <del>-</del> -	-	•	_	Ψ-	•-	-	_	_	_	_	-	-	•		
0.82	0.26	1.28	9	5.4	433	3 5	3.82	70.5	. <del>.</del>	2 Y	, E	? <del>.</del>	7 7	5 4	6.0	6.7	2 5	9.7	7.7	1.1	2.13	0.90	1.51	1.80	3.81	1,40	0.55	3.19	60.	2.65	9	2.05	0.95	0.83	1.12	3.42	2.81	1.83	2.19	18.	46	4.	5 t. t	17.0	0.00	9.5		
0.70	0.73	10.0	67.4	9.0 8.0	2.28	2.78	9 to	0.97	9	C4.4	8 5	2.07	0.30	66.0	4.59	53	0.75	0.68	1.58	1.78	0.78	0.00	1.24	1.12	13		242		77	) ·	) (i	 	5.5	 	9 6	9.00		10.0	9.5	<u>ئ</u>	<b>5</b> 9	05.7	0.82	1.70	0.77	4.12		
:	7.10	0.09	<del>.</del> 8.	09.0	4.37	1.13	0.31	2.69	4.22	0.98	1.51	0.80	1.61	2.17	0.55	0.71	<b>2</b> .	0.67	0.55	1.14	3	233	4.34	7.6	60.0	0.31	5.03	1.98	07.	2.18	1.40	4.79	1.13	1.18	99.0	7.7	1.2/	0.86	1.29	0.86	2.08	1.67	0.48	0.49	0.95	1.76		
,	96.0	0.10	1.42	1.42	1.12	2.24	1.48	1.33	2.98	1.40	1.50	1.41	1.40	1.29	2 29	1.53	2.76	80.0	8.0		9.6	4.24	1.91	27	1.80	1.58	0.65	4.68	2.37	<u>+</u>	1.10	2.47	3.34	1.83	1.03	2.36	1.24	1.07	54.	2.10	8	1.65	98.	28	2	0.75	3	
	0.88	9.46	2.07	20.3	2 7 7	5 5	8 8	1 64	2 2 2	2.45		- c	7 00	D 10 1	6.6	57.7	D (	79.4	6.30	5.11	1.39	1.18	4.21	0.98	2.47	2.88	1,78	2.54	0.73	40.4	3.43	0.68	2.57	4.75	6.75	1,57	1.63	3.84	494	92.6	2 22	1 6		- C	67.7	5.63	/c.L	
	,	- 2	2.10	2.10	2.10	2.09	2.03 00 00 00 00 00	2.09	2.09	2.03	2.07	2.07	2.07	2.07	2.07	5.06	5.08	2.06	2.06	2.06	2.06	2.05	2.05	2.05	205	2.5	2	5 5	5 6	3.5	5.63	3.5	2.63	2.63	3 5	20.5	20.5	7. C	4.04	7.07	70.7	2.02	2.02	2.02	2.01	2.01	2.01	
		0.57	1.28	96.0	98.0	1.76	3	0.01	0.40	0.45	1.18	0.24	1.01	0.64	0.41	2.43	0.47	0.24	1.73	, r.	70.0	2 53	7 70	8t. c	9 6	7 F	c :	1.40	0.60	64.78	0.65	0.28	0.38	1.82	0.43	6.0	F	0.4 0.4	0.62	0.10	3.42	1.06	0.03	5.26	1.73	1.07	2.71	
		1:21	2.72	2.02	1.80	3.68	0.09	0.02	0.83	0.95	2.46	0.50	2.08	1.32	0.86	5.02	86.0	670	7.57	9	9 0	9 9	9.10	23.68	0.53	<del>.</del> 8	15.75	2.86	12	130.51	1.32	0.57	0.77	3.68	0.87	1.97	3.86	0.81	1.24	0.19	6.91	2.14	0.07	<b>10.60</b>	3.49	2.16	5.46	
		N63567	AA935560	W96134	T56726	AA485352	AA292429	W72749	AA757717	AA699573	W96155	H28710	AA150918	N62403	H45447	172915	HOSAAS	D31701	A A 22 A DA 5	AA775.18	2000	A4443502	AA700054	AA777670	AA677050	AA015959	AA996131	H26426	T54164	<b>T64469</b>	R51535	T99852	AA481311	AA463635	AA057156	N48050	H93381	H04382	T61050	AA071503	AA521247	H08424	AA449762	H63416	164223	H93335	A1004169	
		278242	155652B	358531	20000	91008	714498	346119	396098	437481	168631	40665	FORDED	00000	40000	103436	1710	2004	3/5	759948	378461	771274	435036	448885	454589	360684	1603583	162199	68988	80484	38691	123225	796313	811827	489079	281659	241821	149373	76988	366039	827152	45823	785816	20000	80221	342064	1631807	
		263	3 3	<b>5</b> 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7 5 6 7	8 8	8 5	107	997	807	27.0	; £	7 [	3 2	¥17	9/2	9 1	717	278	278	280	281	282	283	<b>784</b>	285	286	287	288	289	8	2 5	292	293	35	282	296	297	208	660	Ş	3 5	3 5	3 5	3 8	\$ 8	2 2	3 8	Š

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1.49	3.72	, K	2 6	2	- <del>-</del>	33 •	8 6	8 1	Š.	<b>3</b>	0.97	C 32	.0 <u>.</u>	1.14	<del>2</del> 1	5.73	1.03	0.93	4.11	<b>8</b> .	1.41	2.39	4.07	2.14	0.68	0.60	0.88	2.84	2.81	1.37	4.37	1.89	3.02	126	8:5	<b>8</b>	1.36	0.81	3.35	1.63	2.46	2.18	1.50	3.54	0.85		
7			1.42	ò :	8	76.1	1.03	<b>-</b>	4.19	0.75	<b>3</b> .0	1.48	1.53	1.52	<del>.</del> 2	0.53	0.61	0.73	2.22	1.50	0.65	0.81	0 70	2.02	1 0	0.00	4 4	4 12	1 29	1.70	1,33	1.60	1.70	96.0	1.52	0.93	2.79	0.71	2.82	0.93	0.14	1,2	0.79	18.0	280	5	
,	200	0.62	5.26	5.61	1.08	2.15	2.39	4.42	1.08	0.25	5.14	3.80	3.89	2.54	1.29	6.33	0.70	0.99	4.0	1.48	3.74	2.17	: C	9.50		9.07	7. T	4. C	9 6	22.5	140	. e.	0.39	0.85	3.47	2.39	1.27	0.93	1.58	<u>4</u>	0.09	3.68	2 55	÷ 52	7 0	80.0 0	
,	1.27	2.08	1.81	1.71	1.06	3.32	1.35	1.87	1.11	2.62	<u>+</u>	2.31	1.96	1.42	1.82	1.17	1.1	86	1.48	0.85	2.60	98.0	9 6	3.5	5. 6	0.78	9 6	9.0	D .	S	 	5 5	5	1.12	1.49	3.4	0.57	2.15	0.93	184	2.89		3 3	2 <del>.</del>	5 6	0.52	
!	2.43	2.81	0.81	66.0	5.19	1.73	3.60	1.47	2.04	2.98	1.75	1.38	53	3.29	3.41	-	6.42	5.73	1.57	<u> </u>	3 3	<u> </u>	3.50	1.01	1.26	06:0	1.7	1.80	86.0	3.17	 	7.7	0.00		2.21	1.56	3.68	5.07	86.0	3.44		0.7	<b>5</b> ;	٥/١	7.47	6.84	&
	2.01	2.01	2.01	2.01	2.00	2.00	2.00	2 00	66	60	60	60	86.	9 6	0,0	000	1.30	) P	.e	.e.,	1.97	1.96	1.96	1.96	1.96	1.96	1.95	1.95	1.95	1.95	<b>3</b> .	3, 3	<b>5</b> . 3	<b>5</b> -	3	3	5	8 5	501	6	2 5	3 5	£8.	1.93	1.93	1.93	
	4.73	2.33	0.92	063	0.77	0.48	4. 1.	88.	3 3	\$ 5	ر ا ا	9 6	5 c	4 5	2 6	9.7°	9 6	67.5	7.52	2.52	3.21	0.28	6.16	2.46	1.06	1.24	0.01	1.19	2.47	7.09	0.43	2.22	<b>3</b> . 9	5.12	6.53	. · ·	. <del>.</del>	- 6	0.30	7.75	5. Ç	Z	90.0	4.49	9.0 \$	1.58	
	9.52	4.69	1.85	1 24	7	65.0	333	ŗ	? ?	2 5	7.07	9.5	77.0	7.47	7.0	5.70	G (	9.49	2.83	4.57	6.31	9. 5.	12.08	4.83	2.08	2.42	0.02	2.31	4.82	13.82	0.85	4.32	2.02	<b>3</b> 3	\$ \$	2 5	7 5	B .	• F	27.0	9.7g	*	0.12	<b>8</b> .66	1.24	3.05	
	AA708440	H38650	W15263	HALIAG	N38860	AARIJARR	A A 630771	1,000	1,6547	115417	V A 404703	7467583	AA404833	2017133	A4212/U	7600/1	/0809A	AA453015	H1405/	N896/6	AA486435	R42520	H42679	R96827	AA427740	AA018480	AA443649	R34331	AA453997	AA962236	H23137	AA009608	AA128407	AA055835	A70/0/2	767248	14/4443	M00000	124431 44403387	VOSSO / V	W129829	HISTORY	<b>W</b> 111100	AA121836	R08266	N91914	
	506016	190732	122547	240676	270005	0.00	956.420	07070	243003	49555	233464	138300	838899	362628	731031	83338	416556	788334	163841	307053	842918	29954	183337	200741	770837	362424	771258	138474	795262	1602120	52303	365515	526587	37748	413150	429148	80220	410833	132954	450050	548683	50877	449257	568106	127193	306829	
	٤	3 8	3 5		- ÷	4 6	2 3	7 1	6	919	317	318	318	8	22	322	23	Ř	328	326	327	328	328	8	33	33	33	ğ	88	338	337	<b>8</b> 2	88	z	<del>,</del>	75	3	3	£ ;	ž	7	<b>3</b>	3	350	351	352	

	-	- +-		· <b>-</b> -	-	<b>~</b> -	-	-	-	-	-	<b>-</b> - ·	- ,	<b>- •</b>	- •	- •	- +	- +-	. 🕶		-	-	-	-	-	<b>,</b> ,	<b>-</b> 1	- <b>•</b>	- +	- +-	. 🕶	-	<b>.</b> .	- •				. <b>,-</b>			
0.94 3.28 1.21	0.86	7 C	7.47	1.02	1.5.	23.7	÷ +	181	3.83	1.30	2.43	0.65	3.44	1.59	0.46	1.87	1.65	64.	4.1.	0.0	2 5	3 5	1.30	5.67	1.90	2.89	0.78	1.26	1.68	44.	- 5.5 - 5.5	2.52	0.93	1.14	2.41	4.35 5.5	8.5	146	) :		
1.97	1.40	2.30	2.41	1.36	1,36	1.97	3.51	0.78	1.75	2.10	2 4	565	58.0	0.57	0.80	1.48	2.42	1.31	0.87	<del>.</del> 3	1.50	0.87	0.73	<u> </u>	 	- C	1.05	0.75	1.02	1.09	1.07	0.92	1 12	1.52	1.93	202	3.32	0.97	0.60		
5.07	0.58 7.69	86.0 66.0	0.41	45.	1.62	3.12	1.90	1.01	0.67	0.42	<del>.</del> 9	1.41	1.97	0 to	5.0	4 47	- 60	3.16	1.86	5.41	1.87	0.86	99'0	3.74	0.41	1.86 2.	1.28	9 6	2.24	1.7	0.70	5.59	0.31	8 6 7	4.38	66.0	4	3.01	1.58		
0.71	0.82	1.16	<b>5</b> 6	5 / C	7 S	3	700	1.54	90	1.31	1.55	1.56	2.54	2.64	3.82	0.81	2.	1.16	84. 84.	1.00	1.17	5.23 8.00	5 5	1.83	1.18	1.13	1.42	52.	2.04	8 8	99.0	1.27	0.63	1.28	0.89	7.0.4 3.5	3.73	25.55	1.36		
9.0 29.0 0	5.76	3.51	3.37	3.58	2.33	3.35	<b>4</b> (	0.85	44.0	2.29	 	5.02 101	3.70	1.77	2.47	0.71	3.33	1.16	2.02	4.59	1.02	1.69	4.90	5.55	 	60°E	3.00	0.72	4.39	1.72	3.46	¢ 6	2.37	5.12	1.18	3.17	1.16	2.40	1.70	σ	ı
8.	1.93	26	25	1.92	1.92	1.92	1.92	1.92	1.92	1.92	191	1.91	1.91	5. 3	 	S 5	06.	9.1	1.89	1.89	1.89	1.89	1.89	1.89	1.88	1.88	00.7	0°.	1.87	1.87	1.87	1.87	1.87	8.4	8. 6	1.86	1.86	1.85	1.85	3	
1.39	0.81	5.43	# C	) (2)	5.00	91.0	2 6	. o	0.43	0.53	8.71	6.47	3.02	92.35	<b>3</b> .	2 2 3	0.76	2.17	3.5	1.30	7,0	700	1 63	0.80	5.86	0.40	0.80	0.02	0.52	1.2.1	0.30	0.49	33.03	12.25	5.5	.0. .84	2.03 10.0	0.39	0.93	0.5/	
	1,57	2.79	1.80	1.53	7.48	0.03	5.50	98.O	9. C	7 E		12.38	5.78	176.33	22.04	1.2	1.45	1.4	2.52	7 F	7.7	0.50	7 .C		11 03	0.76	1.51	9.	0.98	2.28	97.0	3 6	49.19	22.82	1.88	3.78 5.68	4. c	20.0	1.72	1.08	
1able 2-7	AA883735	AA953973	AA426113	AA235370	AA670286	AA130017	AA460827	AA666418	AA085748	140905	100001	N50912	N91003	AA055768	AA682627	H74086	N26171	AA155695	AA975556	N73301	AA504492	AA776434	9/989N	H09935	A A 3 0 B 0 2 2	R53963	N64391	R05416	H11376	AA088517	AA757659	AA400014	AA873355	AA775447	N92801	AA931725	AA626315	H5Z361	AA707840	R26859	
	192569	1573305	757265	687667	878281	589362	796268	859654	488301	61412	45877	841615	281072	300000	431786	214809	269425	592243	1555451	292068	825356	453766	292633	46358	727792	727164	35833	125134	47630	511832	396045	701272	195127	978187	307660	1572196	745570	202168	32578	133358	
	353	\$ 15 15 15 15 15 15 15 15 15 15 15 15 15 1	3 5	3 5	5	359	360	8	362	88	36	365	98 i	36.	Š	8 6	5,5		37.5	374	375	376	377	378	378	တ္တ	<u> </u>	382	3 3	38	386	387	388	80 8	9 5	302	383	<b>3</b> 6	395	397	•

Table 2-7: Good vs. Poor Clinical Outcome: Genes upregulated 3 fold or more in at least 1 tumor sample.

1.25 3.19 0.97 4.40 0.95 0.86 1.48 1.85 1.03 1.17	2.44 1.74 0.63 0.84 1.44 1.19 1.19 1.37 0.00	2.35 1.54 1.02 1.11 0.99 0.97 1.20 1.20	0.56 5.40 1.09 1.09 1.01 1.73 1.73
2.54 1.45 1.45 1.16 1.52 2.00 0.88 0.89 0.89	1.72 1.67 1.67 1.08 0.80 1.71 1.71 0.88	2.2 2.0 3.0 3.0 2.2 2.2 3.3 3.0 5.0 5.0 5.0 5.0 5.0 5.0 5.0 5.0 5.0 5	1.18 1.28 1.06 1.05 1.20 0.80 1.73 1.73
5.01 6.48 7.17 7.161 7.54 7.54 6.09 7.77 9.77 9.86	3.18 1.68 1.68 1.08 3.05 1.07 1.07 3.33 3.33 4.14	0.66 3.26 4.97 3.19 2.29 4.08 1.99 0.30 3.28 3.14	3.51 0.79 0.79 1.95 3.63 3.73 0.81
1.05 1.17 1.17 1.01 1.25 1.56 1.50 1.72 1.72 1.72 1.72	1.10 1.10 1.22 1.22 1.30 1.30 1.40 1.40 1.65 1.65	1.50 1.76 1.94 0.94 0.98 0.98 0.98	1.05 0.93 0.93 0.77 1.44 1.63 1.76 1.75 1.75
0.78 1.91 1.40 1.02 3.24 1.85 4.95 4.95 6.85 8.95	3.50 0.62 0.43 3.52 1.18 1.10 1.10 1.33 1.33	3.36 1.15 0.80 1.59 1.64 1.76 2.99 2.99	3.20 0.59 3.65 1.57 3.23 3.23 1.35 0.58
28.1. 28.1. 28.1. 28.1. 28.1. 28.1. 28.1. 28.1. 28.1. 28.1. 28.1.	1.83 1.83 1.83 1.83 1.82 1.82 1.82 1.82 1.82	1.82 1.82 1.81 1.81 1.80 1.80 1.80 1.80 1.80 1.80	1.80 1.80 1.80 1.79 1.79 1.79 1.79 1.79 1.78
1.27 15.38 0.87 0.12 1.27 1.47 2.52 0.80 0.56 0.25	0.46 0.11 1.10 1.10 1.189 0.03 0.03 0.03 37.83	2.47 2.47 2.47 2.64 2.26 2.26 2.95 15.33	0.60 0.39 0.39 0.51 0.45 0.63 14.15 18.60 10.58
2.36 28.41 1.61 0.21 2.35 1.19 2.71 4.64 1.47 1.03	0.85 0.21 1.30 2.00 2.00 0.56 0.06 0.08 0.90	66.74 4.49 1.58 13.83 16.82 0.46 2.43 4.62 1.72 2.762	1.08 0.70 0.70 0.92 0.92 1.13 25.27 33.23 15.09
AA704587 AA886870 H22563 AA490987 H19826 AA699714 AA43188 W32408 R80217 AA505045 H17896	A441337 A441933 AA688516 AA689649 H20547 AA019482 AA019482 H10761 R48320 AA707835 AA086886 N47664	AA772833 AA872454 AA199881 H98075 AA55042 AA975209 AA78223 AA521394 AA521394 AA57354	K97226 K48833 AA882293 AA882184 AA898184 H08747 N32904 AA609474 N35067 H23211
450598 1493252 61916 824573 172517 433287 796876 321359 147050 825648 60043	754601 774078 1408710 1031278 51320 489706 383086 47169 153541 392399 489553 2778448	87854 1476019 64528 262080 81224 1555427 809620 611412 808038 828072	200395 163617 461727 701112 1580342 46353 258684 1031598 271662 51773
388 400 401 402 404 404 406 608	4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4	25	555555555555

0.50	98.0	8 6	9.6	20 ·	1.57	2.65	0.77	3.76	3.43	4.42	308	0.65	3 5	3 5	5 3	5.5	C	9 9	0.63	35.	5.76	1.07	0.56	1.05	3.58	1.52	0.68	1.26	1 23	0.85	60 4	26.0	8 6		08.0	2.00		2. 4	27.	3 6	0.30	0.70	0.63		0.81	4:-	1.43		
7.77	57.4	<b>5</b>	<del>-</del> :	1.28	1.65	3.22	0.89	1.30	1,95	1.02	10.		<b>5</b>	0.80	18.0	1.22	0.56		2.46	1.74	1.52	<del>1</del> .	1.37	1.71	2.07	3.28	0.81	1 65		6 6	 		2 7	<u>*</u> ;	0.32	28.0	9.5	 6. t	7.7	3.C	00.0	<u> </u>	1.40	0.71	3.61	3.32	3.49		
90	8 8	3.02	0.80	9:	1.30	0.84	1.29	0.21	0.38	+ + + + + + + + + + + + + + + + + + + +	1 5		4.4	1.7	3.74	1.68	3.20	0.83	3.15	3.32	0.35	3,68	4.08	3,45	0.27	5	25.4	4.44	- c	0.0	9 9 9 8	0.32	6.29	0.48	9.00	0.7	3.61	3.01	0.89	1.07	4.19	3.01	3.80	3.99	96.0	1.37	1.38		
į	1.72	1.90	4.45	1.39	0.87	1.40	4.71		8 -		70.7	1.12	0.85	<b>2</b> .	1.98	1.85	2.20	2.	1.07	1.15	0.61	0.82	181	1.45	8		2 7	<b>*</b> 00	0.80	ς ο :	80	1.29	0.79	<b>5</b>	E	2.66	2.01	1.89	0.89	1.29	2.01	1.15	1.06	2.12	2.29	0.98	1.68		
	1.42	2.15	1.44	3.75	3,50	0.74	5 5	32.		<u> </u>	0.47	1.32	÷.90	3.44	1.18	3.05	2.31	2.13	1.47	1.02	0.55	7 · ·	- <del>-</del>		* · · ·	0.93	1.78	1.28	3.54	4.98	1.52	2.31	3.28	2.26	0.78	3.59	1.16	1.33	3.53	1.45	1.00	1.27	1.55	1.13	0.98	1.85	0.66	•	=
	1.78	1.78	1.78	1.78	1 78	F	<u> </u>	<u> </u>	<u> </u>	<i>!"</i> !	1.71	1.71	1.7	1.77	1.76	1.76	1.76	1.76	1.76	1.76	2. t	1.70	). 	D	C/.	£ ;	1.75	1.75	1.75	1.75	1.75	1.75	1.75	1.74	1.74	1.74	1.74	1.74	1.74	1.74	5.	1.73	1.73	1.73	1.73	1 73	, r	<u>:</u>	
	4.41	1.49	75.0	2	80.	8 8	3 6	8.0 8.0	5.95	20.	5.60	3.15	0.21	0.95	0.83	0.82	67 79	0.40	600	5 6		S : 5	6.5	/c:0	0.11	11.60	2.77	0.89	1.23	1.12	1.41	1.06	0.57	0.97	0.91	0.91	8.20	0.07	0.90	1.50	0.61	0.57	1.16	1.42	1.12	33	30.0	20.0	
	7.87	2 65	3 5	5	- F	3.	3.5	0.68	10.53	2.73	9.91	5.57	0.37	99	1.47	45	118.87	2 2	, 4	5 6	5 5	2 6	<b>S</b>	90:	0.18 8	20.33	86	1.56	2.16	1.97	2.46	<u>~</u>	<b>5</b> .	1.69	1.59	1.58	14.29	0.12	<del>.</del> 8	2.60	8	0.98	200	2.46	5 5	. c	3 8		
	AA442853	AA485140	AA478479	MEAROR	A 4 20003	CERO35	Aidodouz	AA707336	R53455	T66320	T97762	W69271	H19217	N51069	AA086204	N78092	AA453310	AA777394	A A 775774	DECREA	100001	CARRON	AA460849	AA652452	AA629804	AA152347	N47902	W49583	AA099523	W88737	N34395	AA459123	N39426	AA283007	H58702	AA278757	AA426025	AA205413	AA455497	AA989210	N20482	H52247	AA456569	AA479351	44496359	AAR7MSR	A 4 50 4 5 4	5 601 5	
	757873		70101	00007	245330	7.77(2)	1809036	451511	39833	21894	121948	343760	51103	281936	401333	248288	700100	700100	448033	67023	138294	295843	786287	450612	884539	504791	281442	325001	489594	417777	277266	814341	276871	713129	206288	703577	757244	646657	809719	1601852	284188	235909	809374	753907	755200	070744	1 40/0	608835	
	443	?	<b>!</b> :	7 :	<b>9</b> !	Ž	<b>\$</b>	<b>4</b>	<b>장</b>	451	25	3	Ž	£ ₹	3 5	3 5	<b>2</b> 4	3 4	g ç	<b>3</b> 3	ē i	<del>2</del>	<b>§</b>	204	465	<b>4</b> 68	467	<b>8</b> 9	469	67	471	472	473	474	475	476	114	478	479	480	481	£ 2	8	787	Š	000	<b>§</b> :	487	

-	• •	- ,		-	-	-	-	-	-	-	-	-	•		-	-	-	_	-	-	-	-	•		•				-	· •-	-	-	-	-		-	-	-	-	-	-	_	-	-	<u>.</u>		
9	<u></u>	1.24	3.23	3.05	4.02		0.53	0.97	1,43	0.98	76.0	5.6	9 6	60.5 F	3 5	5 6	7	990	23.5	76	5.5	,c.,	1		2 .	07.5	7 7	1.37	. c	670	1.32	1.19	3.68	1.29	0.80	3.68	0.91	2.46	1.41	0.82	3.73	2.00	0.42	3.42	2.25	ì	
;	3.	1.59	- - - - -	2.38	1.07	52	0.28	790	9	22.5	÷	2 3	5 7	1.61	5.23	8 2		8. 5	- c	- c € £	S	<b>*</b> 8	5 G	8 8	8 8	<u>8</u>		20.1	D	\$ C.	5.4	2 6	1.57	3,55	66.0	22	96'0	1.37	1.80	1.26	122	1.18	24.5	25	5	•	
	3.64	<b>3</b> .0	0.93	1.74	500	200	7 6		3 5	n 1	CC.1	4.28	1.15	1.46	1.35	0.50	07.	3.25	3.79	0.65	0.58	3.05	0.41	1.27	1.43	0.21	0.48	3.10	3.01	1.05	9.00	9 6	. c	203	4	4	5.10	0.44	1.45	3.38	9 6	176	70	5.6	56.0		•
	1.27	1.00	35	0.78	2 6	20.0	70.0	<b>7</b> .7	20.6	1.12	2.35		1.80	1.32	1.48	5.09	1.52	1.17	1.38	0.88	0.58	0.91	1.23	1.36	0.71	1.20	1.12	1.23	1.16	0.85	1.77	1.24	. c. 33	\$ 5	28.0	2.0	690	0.83	9 6	8.5	7. 4	- 6	0.0	2.5	1.18	0.43	
	1.21	3 05	4.06	3.0	000	1.6/	1.84	4.03	0.71	1.27	3.09	0.84	1.27	3.25	1.19	1.46	3.97	1.20	1.78	4.26	3.03	1.82	3.67	4.10	3.79	2.63	3.32	1.46	7	29.0	1.42	0.68	0.70	- 13 - 13	8 5	\$ F	3 6	37.0	0.40	- ;	- 1	9 70	98 (S	0.71	1.48	3.15	12
	1.72		7/.	7/1	1.72	1.72	1.72	1.71	1.71	1.71	1.71	1.71	1.71	1.71	1.71	1.71	1.70	1.70	1.70	1.70	1.70	1,70	1.70	1.70	1.70	1.70	1.70	1.70	1.69	1.69	1.69	1.69	1.69	1.68	80.	89:	8 5	70.7	) o'.	7.67	1.67	1.67	1.67	1.67	1.67	1.68	
	75 6	£.5.	E.4.	0.30	0.37	1.30	2.28	3.16	30	1.89	0.42	09'0	78 0	1.43	2.53	0.40	1.89	2.28	0.42	2	19.38	4 95	3.40	1.89	99	15.91	1.40	0.59	161	1.02	2.	0.37	0.30	29.16	1.18	2.85.	8.50	3.02	S	17.98	0.82	1.26	6.59	<b>7</b> -0	0.88	2.08	
	2	90.1	7.78	0.52	<u>7</u>	2.24	3.91	5.42	0.07	3.23	0.73	5	<u> </u>		2	0.78	3.22	3.88	0.72	43	32.62	3 T	4	2 5		20.5 20.5 20.5 20.5	3 %	8 8	27.0	<u> </u>	2.77	0.63	0.50	49.12	1.99	4.78	4.20	5.05	7.21	30.10	1.37	2.11	<b>\$</b>	0.73	1.46	3,46	
	4 4 704 127	W.29113/	AA598796	AA447514	R51617	N2634	N40951	R31413	AA156873	AA476240	W39609	AA454562	AA285053	T47625	H24316	AA018215	R00128	R07184	AA706870	AA504845	A A 418877	A4782292	A A D.4.2000	AA609731	AA400317	AA488986	B01515	AA678087	AA405800	R06716	AA496348	W70343	R42736	AA505141	R25403	R39446	AA777355	AA196485	H28091	AA181179	R53943	W86282	AA019335	W23631	AA084517	AA416767	
	6	700503	860868	782576	39442	285668	277185	135065	502527	771323	322786	25.00	11111	71317	11312	28720	423061	426879	411482	43143	721200	087157	00000	186034	2355	743533	67/470	078571	77777	128474	755274	345849	31366	825822	133179	23819	448417	628338	162310	624271	40027	415851	363081	327799	546600	731311	
	•	488	489	<b>6</b>	491	492	493	9	405	<b>4</b> 9 <b>4</b>	407		0 0	<b>3</b> 5 5	2 2	2 5	3 5	3 3	<b>\$ \$</b>	2 2	8 8	2 5	0 0	3	2 :	<u>.</u> :	710	25	* 1	U 4	2 5	518	519	520	521	223	523	524	525	526	527	528	529	230	531	23	; !

Table 2-7: Good vs. Poor Clinical Outcome: Genes upregulated 3 fold or more in at least 1 tumor sample.

		<b>,</b>	-	-		-	-	-		- •	- •	- •	- •	- ,	- •		<b>-</b> ·	<b>-</b> ·	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	<del>-</del> 1	-		-		•	-	_			
4.29	1.15 5F.0	25.5	- C	4.16	3.52	46.5	5 6	- 6	9.6	3.52	51.1	0.56	0.92	0.51	3.60	3.98	2.08	1.33	0.83	1.51	1.51	0.71	0.89	131	0.95	7.0	0.70	68.0	0.65	3.69	0.61	1.32	0.82	1.4	0.99	1.29	1.08	0.56	1.02	3.07	1.76	1.27	0.98	9.3 4.0			
1.30	3.51	0.76	0.90	0.0 6.0 6.0 7.0	7 1	270	0.71	2.35	3.61	2.14	0.93	0.49	1.33	0.46	1.66	0.20	0.77	0.65	1.62	8	78.0	1 19	2 5	5 5	3.5	9.0	5.50 9.60	9.60	5 F	 	. ·	2 5	25.0	1.25	1.1	1.20	1.40	0.97	0.68	0.46	1.1	1 29	1.24	848	9		
0.24	2.04	0.50	1.48	4.56	0.70	0.87	0.70	0.79	0.89	0.32	2.59	6.16	1.40	1.20	75	180	1 25	27.0	4 6.73	- + - +	 	2.45	0/.	3.24	1.36	1.30	2.07	3.27	3.03	3.29	0.51	4.65	3.3 3.3	70.7	6 c	3.43	3.58	3.07	3.47	- F	77	1 6	0.63	5	4.0		
0.70	1.02	2.97	3.47	1.53	1.35	2.02	46.0	0.67	780	1 07	, c	3 6	0.00	<u> </u>	. 17.7	5.6	4.7	5.13	1.26	0.70	1.53	2.33	0.95	0.80	1.32	3.63	96.0	1.17	2.11	1.78	1.86	0.86	1.25	3.33	0.48	- 6	0.6	0.00	/8:0 •	3 3		<del>.</del> 5	1.24	98.0	6.24		
7	97.0	933	1 15	0.67	0.67	1 12	- 6	5 F	9 4	6.0	0.10	0.5/	0.46	3.14	0.36	9.	0.31	0.91	4.25	3.97	3.03	3.08	3.59	1.23	3.21	1.35	1.10	2.06	1.12	1.68	0.62	0.83	1.20	0.67	4.08	0.93	1.19	1.18	2.53	1.92	1.60	3.30	3.38	1.41	0.54	13	
	1.66	0.1 99.	99.	8 %	8.5	99.	8 9	8.	99:	1.66	1.65	1.65	1.65	1.65	1.65	1.65	1.65	1,65	2	1.64	2	40.	19	2. 3	5 4	5 5	 		9. 5	6.5	8.5	3.5	163	3	1.62	1.62	1.62	1.62	1.62	1.62	1.61	1.61	1.61	1.61	1.61		
	13.12	2.10	0.60	0.40	PG:0	0.58	26.77	1.38	2.14	1.42	20.37	0.73	0.76	1.50	2.07	40	0.6	2 03	800	80.0	9		9 6	8 2	40.0	81.0	4.20	3.28	<del>1</del> .33	0.48	2.0	3.20	2.0		4.14	1.14	1.59	4.82	0.05	0.40	0.08	3.39	24.		. 60	) } :	
	21.82	3.50	0.0	0.67	86.0	0.97	44.42	2.31	3.55	2.35	33.63	121	1.25	2.47	34.	, e	18 25	<u> </u>			7.57	9. 6.	2.39	3.38	5.81	0.30	6.83	5.37	2.18	0.78	0.19	5.20	0.57	97.	3 5	285	2.57	7.87	30.0 0.09	790	10			5 .	<u>*</u>	,	
	T59658	R77512	AA459110	AA402760	R38678	R52796	AA872257	N77263	AA115742	AA429034	AA487215	T52840	H08194	A A 459293	AA001841	140377	A A C 20117	AA02311	AA/0232/	AA469213	/8005X	N63/2/	AA644099	174023	AA630445	AA599741	R53951	AA464163	AA488976	N52837	AA480870	AA486669	AA018320	N39808	A 6 78 8970	A A 578075	N50843	A 4505082	AA46866	A A 7 M 3 7 1 G	A1017240	A1017.40	1462366	00025W	747,000	190964	
	78169	145491	814320	727026	23443	41648	1477479	245824	400647	760708	007.70	99130	68091	40391	108018	42/390	50926	1035765	447404	825060	137275	293539	845380	84560	854633	1070062	40136	810358	824720	244391	814547	840990	363055	288959	1048789	1240220	383528	00000	872209	01260	788654	1636251	782446	325526	325111	109316	
	613	3 5	35	8	537	<b>5</b>	3 5	3 3	ξ :	<u>,</u>	<b>7</b> .	3	3 3	χ. Σ.	546	\$	25 85	35	220	551	552	553	525	555	558	557	228	559	260		282	\$83	쟗	565	286	267	20 20 20 20 20 20 20 20 20 20 20 20 20 2	8	570	27	212	ટ્ડ	274	575	276	211	

Table 2-7: Good vs. Poor Clinical Outcome: Genes upregulated 3 fold or more in at least 1 tumor sample.

-	-	-	-	-	-	-	-	-	•	• •	- •	- •	- •					•			- •	- •					- •	- •				-	-	-	-	-	-	-	-	-	-	-	-	-		
0.84	Ξ	1.24	3.71	0.78	55	0.88	171	, t	2 5	3. 4.	5. L	CC.1	3.2/	0.73	5.43	S 5	2 85		7:77	ر ان	3 6	96.0	8 5	3 5	¥.07	0.70	2.5	5 4	7.10	78.0	<b>t</b> 95	89	5	1.03	0.97	0.63	3.31	1.09	3.29	<b>5</b> .	0.82	0.60	2.13	3.26		
1.23	1.06	3.08	0.87	8	1.47	1.46	7. 4		0.1	<del>.</del>	2.13	111	2.23	1.70	2.0	0.00	<b>X</b> :	\$ ;	1.28	<del>.</del>	7.00	1.19	0.76	1.01	1.36	3.26	1.17	(6.9 (0.9)	S F	7.0	2.24	6 ° °	1.33	0.72	26.0	3.20	1.98	0.99	2.24	1.01	3.79	1,11	0.84	1.60		
1.05	1 10	22.6	950	900	60.1	5.50	9 9	. C	3.05	<del>.</del> .	0.93	0.78	0:30	3.47	3.54	3.18	3.80	3.4/	<b>1</b>	0.65	3.68	1.10	1.00	0.97	0.22	1.68	4.39	1.05	3.42	3.21	1.03	0.50 0.50 0.50 0.50 0.50 0.50 0.50 0.50	3.04	. c. c	980	1.06	0.33	3.22	0.50	3.55	1.95	1.20	0.76	1.27		
101	35		7 6	D 7.1	3.17	0.62	J. 1	0.47	1.08	0.78	0.73	3.29	2.	0.67	2.31	1.46	2.82	1.05	0.49	1.35	1.75	1.08	1.28	1.24	0.78	1.09	1.13	<u>.</u> 8	1.7	0.75	0.60	88.68	£ 5	8/:O	S &	- c	2 2	130	290	1.12	8.	<b>7</b> 80	0.50	0.87	į	
, 89	66	B .	<u> </u>	<b>5</b>	2.11	<b>*</b> .03	0.64	1.10	1.29	1.22	<b>.</b> <b>.</b>	1.44	1.12	1.30	0.90	1.02	0.52	1.16	3.17	3.56	0.82	3.51	3.97	3.42	1.43	1.05	0.38	3.17	0.59	1.36	3.13	0.87	3.66	0.50	3.78	3.72	1.35		9 9	1.05	0.12	3.99	3.52	0.73	14	_
9	8.	9:	1.60	1.60	1.60	1.60	1.60	1.60	9.1	1.60	1.60	1,59	65.	1.59	1.59	1.58	1.58	1.58	1.58	1.58	1.58	1.57	1.57	1.57	1.57	1.57	1.57	1.57	1.57	1.57	1.57	1.56	5.5 8. j	<del>.</del> .	92.	8 5	0.7	8. 4	3 3	3 5	3	- 	3 4	3 4	3	
ç	स्तु । •	0.53	<b>4</b> 8	2.15	0.75	1.74	.0 <del>0</del>	2.8	0.51	6.26	4.46	181	3.47	27.4	\$	0.29	0.99	1.75	1.37	0.43	0.72	00 6	0.34	0.30	8.88	1.	1.85	\$50	6.24	0.47	1.47	3.03	0:30	15.46	0.55	0.7	0.60	6.7.3	60.0	2.10	2.70	20	2 76	2 6	<b>5</b>	
Ş	0.62	0.86	7.28	3.45	1.20	2.78	1.74	11.24	0.82	86.6	7 12	88.	3 4	7.51	2.13	0.45	55.	2.77	2.17	990	1 14	7.2	920	0.48	13.96	1.75	2.91	0.84	9.80	0.74	2.30	4.72	0.47	24.12	0.85	1.10	1.24	67.4	9.5	. <del>.</del>	5 7	2	, e	3 6	6.5	
	R9767H	T99011	AA521431	R63106	R37619	AA479967	N21338	A001449	N24538	AA411686	T55728	AA187207	AA056232	AA434092	AA677534	AA021628	R52901	R24506	R26960	H74265	AA443832	AA663123	R38962	R91503	AA504265	AA399223	R74478	N58170	AA454634	AA214031	W60845	AA677287	AA705319	N47240	AA282134	AA303051	AA778633	77320Z	A A 026474	038047	AAR20715	B44447	AAATESSE	UDBER7	2000	
!	49842	122684	826173	137940	137254	753982	265103	361974	267541	75337B	7.638	624490	084470	101710	460403	364083	41569	131791	133273	22026	784065	657548	24855	108387	825416	725745	143450	247660	811880	562395	341805	454771	462856	280459	711918	825654	1049009	52078	40567	7691/4	101030	7484.50	800 <b>%</b>	5000	230399	
	578	<b>2</b> 20	88	581	582	583	28	585	88	587			8 5	90	9 2	503	3	70,	3 8	201	804			3 5	3 5	3 8	3 8	55	88	607	808	009	610	611	612	613	614	615	919	75	0 0	<b>A</b> C	200		7	

Table 2-7: Good vs. Poor Clinical Outcome: Genes upregulated 3 fold or more in at least 1 tumor sample.

	- ▼	- •	- •	- +	- +		_	-	-	-	-	-	-	~	-		•	-	-	-	-	_	-	-	-	-	-	-	-	-	-	-	-	-	-	-	,	<b>,</b>	- •	- \	- ,	- 1	<del>-</del>		
0.97	1.40	0.93	12.	) ()	3.31	28.0	<b>3</b> -	2.0	28	0.76	95	86.0	0.93	3,85	0.67	0.91	0.95	0.33	77.0	1 20	9 6	66.0	7.7	55.5	1 10	0.76	1.12	1.47	3.30	1.07	1.13	3,46	0.90	1.75	0.74	1.21	1.27	0.79	0.98	99.0	0.78	1.26	1.1		
1.20	1.51	0.95	1.27	<b>-:</b>	1.73	1.47	1.24	1.16	3 !	1.4/	\. \.	1.50	, o	5.00	2 5	c	9 6	3 6	4.0	 	3 5	7.0	67.1	1.03	5.50	1.17	9.50 •	ر ا ا		9	25.5	124	98.0	0.70	5.02	4.0	0.96	1.61	1.52	1.49	1.32	1.20	96.0		
3.23	3.32	4.06	0.81	1.13	0.43	0.68	1.10	3.39	3.19	3,62	3.01	0.88	0.88	3.27	0.45	1.12	3.18	0.42	1.25	<del>1</del> .06	1.37	1.03	1.43	0.39	0.38	3.88	1.27	3.16	0.75	9.7	0.67	88.0	0.20		25.0	- &	20.6	3.20	908	96	3.58	0.81	0.76		
•	1.4.1	2 0	1.27	6	1.1	0.53	1.24	1.01	1.68	0.72	<u>5</u>	1.10	1.75	0.85	12	1.37	1.39	0.70	4.17	1.03	3.56	1.19	0.74	1.25	1.59	0.80	1.71	1.42	1.3 8.	121	0.82	0.93	96.0	28.	39.0	8 6	2.5	 	2.0	5 6	9 6	, c. c.	1 17	<b>:</b>	
•	1.12	3.6	6.03	5.5	5.07	 	3.18	1.10	68.0	0.58	1.21	3.35	3.19	1.53	0.93	3.16	1.26	4.69	4.	3.42	45.0	3.72	3.20	3.15	0.67	0.63	3.22	0.52	3.37	0.59	4.33	4.05	1.64	7.0	3.61	0.48	4.08	0.82	1.10	0.99	69.0	0.82	3.40	3.4.	15
	1.55	<u>.</u> .5	4.	٠. يخ	<b>7</b> 5.	<b>X</b> 3	4. T	đ.	<u> </u>	 	5.4	56.4	5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5	3 5		52	1.52	52	5.5	55.	25.1	1.34	5.35	7.07	4 5	1.52	5.5	1.54	<u>.</u>	, <u>r</u>	55	1.51	15.	1.51	1.51	1.51	1.50	1.50	1.50	1.50	1.50	1.50	1.50	1.50	
	08.0	40.	4.21	0.49	0.23	2.16	2.14	0.10	29.6	0.69	75.80	0.55 0		5.5 5.5	- 5	2.5	0.50		χ. Α	2.30	0.62	9.11	0.88	2.12	2.53	50.4	24.03	0.00	0.5	97.	 	2.0	2	2.25	0.78	0.35	1.76	1.92	1.46	0.75	3.21	0.47	1.20	0.71	
	1,3	8	6.49	0.75	0.36	3.32	3.30	0.29	8.63	1.07	116.27	0.85	<u>6</u>	0.51	2.5	15.46		51.1	2.35	3.59	9.0	13.85	<u> </u>	3.23	8. 4	6.20	37.32	9.78	12.33		 	97.0		3.40	1 4	120	2,65	2.88	2.20	1.12	4.80	0.71	1.79	<del>2</del> .	
	AA598809	R86847	AA489016	AA135886	H15695	AA040617	AA879437	R51912	N51335	AA418414	N52178	N75356	H72959	AA015793	N26724	AA1429ZZ	R82733	AA186405	AA872397	R10378	R33103	AA701502	AA129552	H57857	H02307	AA290737	N48809	R16245	AA159578	T60149	<b>A</b> 733317	AA284113	74406	6042455	NOTATION	NAGARO	AA775810	AA778320	T63031	AA707008	W46985	AA421284	W31725	R44985	
٠	300200	02/370	105040	22520	49203	486208	1505692	39593	283142	767284	284247	298862	213917	360478	266300	505225	149742	628295	1472743	-₫	006000	435470	564803	205497	151201	713922	279481	53393	592594	81408	399421	324751	27769	84613	76/1/0	307231	8476/7	170/9	703808	454354	900000	724054	320797	74442	!
	<b>§</b>	3 3	\$ 8	3 5	97.	628	629	630	5	83	8	<b>5</b> 5	635	636	<b>637</b>	638	639	<b>₹</b>	2	26	<b>6</b> 43	3	55	8	24	848	<b>6</b> 29	650	53	652	653	654	655	656	657	658	8	3	<b>§</b> 8	8 8	3 3	8	2 8 2 8 3 8	8 4	ŝ

Table 2-7: Good vs. Poor Clinical Outcome: Genes upregulated 3 fold or more in at least 1 tumor sample.

<del>-</del>	<b></b> ,	-	- ,	<b>-</b> ,	•	- ,						• •	. —	-		-	•	•	-	•		-		-	· <b>-</b>	-	. 🕶	-	<b>,</b>	-	-	-	_	-			- <b>-</b>	- •	- •	- ,		<b>-</b> ,	-	
4.43	0.82	0.83	1.30	0.52	1.17	2.00	1.65	A	9 6	0 0		<del>?</del> <del>?</del>	10	7 1 7	16 C	5		700	; <del>-</del>		- F	3.25	67.C	2.5	. 60	75	- 0 86	08.0	1.15	1.16	0.80	60 -	3.41	0.87	6.6	9 9	9.6	7.0 9.0	6.60	\$ : :	0.92	0.77	0.63	
0.76	1.75	1,35	1.72	0.85	3.20	0.46	0.63	2.09	1.07	<del>2</del> 6	0.93	57.1	9 5	9 9	0.0		3.43	3 5	2. C	5.6	4.0	07.1	0.80	G 60	0.0 7.8	2,0	0.70 90.0	06.0	10.1	÷ 5	0.67	900	9 6	3	8.5	7.75	8 8	0.65	 90.	2.09	1.37	1.72	0.71	
0.50	3.40	3.49	3.49	3.53	1.15	0.42	0.52	3.00	0.95	3.09	0.98	0.91	9. G	٠ ٢	0.97	25.5	70°C	C1.7	3.16	3.27	0.35	5.03	00.0	<b>6</b> (0)	 	77.4	2. c	/8.0 E	7/5	5 6	8 2	5 6	- 60 C	3 8	3.33	3.23	3.73	3.01	3.93	3.14	0.89	3.04	3.87	
1.01	1,05	1.25	0.58	<del>.</del> .	1.31	3.64	1.45	0.95	1.0	1.45	26.0	0.93	0.73	0.92	1.06	16.0	0.78	3.02	1.05	1.02	1.40	0.97	4 .	1.15	0.79	0.81	1.12	99.0	1,45	5	 6. £	5.5	0.65	<b>5</b>	1.35	0.86	0.82	1.28	0.67	0.69	1.06	0.64	1.08	
0.77	3 7	0.55	0.37	121	0.63	0.87	3.20	0.20	3.51	0.57	3.68	3.07	0.56	1.13	1.51	1.01	0.77	0.85	0.90	1.06	3.96	0.94	1.47	1.24	1.14	0.30	3.49	3.61	3.33	3.07	3.02	3.96	0.62	0.69	0.74	1.12	0.58	1.39	0.65	69.0	3.01	1.06	0.94	16
1.49	. t	7 7	64.	1 40	149	1.49	64.	1.49	1.49	1.49	1.48	1.48	1.48	1.48	1.48	1.48	1.48	1.48	1.47	1.47	1.47	1.47	1.47	1.47	1.47	1.46	1.46	1,46	1.46	1.46	1.46	1.46	1,45	1.45	1.45	1.45	1.45	1,45	1.45	1.45	1.45	1.45	1.45	<b>?</b>
3.07	0.70	5 6	2 ° °		 	77.	0.82	6.83	0.53	0.41	4.56	0.73	15.21	96.0	9.9 75.0	6.02	0.60	1.35	1.25	0.80	0.46	1.40	6,52	5.09	0.48	0.79	0.59	0.49	0.70	0.39	0. 83	0. 51.	24.04	5.67	0.38	4.70	92.77	0.55	2.27	4 05	13	7.46	105	***
4.59	<u>-</u>	<u>.</u>	2.40	3 5	. c.		2.13	10 18	0.78	09.0	6.76	1.09	25.52	1.46	0.50	8.90	0.74	1.99	<b>7</b> .	1,18	0.68	2.06	9.58	7.47	0.67	1.15	98.0	0.72	1.03	0.57	0.92	0.22	まる	8.23	0.57	6.82	134.60	0.80	330	1.52	8	10.78	1.51	<u>.</u>
W16832	AA598817	000000	W8/825		PO8121	A A BTTAKTE	AA131406	R26785	N33443	R44544	AA400457	AA609392	AA702077	R12385	H03436	N69528	AA489729	AA017468	AA962119	AA701900	N58417	AA707400	H08899	N30285	AAB44211	AA678203	AA283949	W86214	R26444	H19320	AA405569	AA453085	H84657	T88721	R52526	AA701046	H62839	AA213669	R93591	R86082	AA233070	AA488064	R28206	
320392	897956	272529	417081	1,0004	614738	127.120	747277	432836	270750	2607	753162	743481	384224	128165	150897	294503	823663	361291	1600281	435536	248117	451546	44975	257504	84547	430864	700571	415868	133150	50918	772425	789069	219888	109863	38977	397432	208377	683278	107731	440474	666492	840A86	433053	3
899	699	670	671	7/0	673	* 10	678 878	3 5	. 44	9 6		681	682	683	<b>2</b> 6	685	686	687	688	689	630	5	692	693	4	695	969	697	898	669	200	5	702	50	ğ	705	902	707	2	3 5	2 5	2 5	= ;	7

Table 2-7: Good vs. Poor Clinical Outcome: Genes upregulated 3 fold or more in at least 1 tumor sample.

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	277	, or r	5 6	68.5	0.78	2.63	22	3.47	. 6	3 1	0.87	0.98	3.05	4.02	0.48	45	5	3 ?	4 8	67.0	1.20	3.5 5.	0.24	1.48	26.0	0.81	<u>5</u>	0.33	8	0.58	1.18	44.	9	33.0	3.13	0.46	0.82	0.82	0.82	16:0	52,	57.0	1.2	1.20	8	80	1.13	5	!		
	;	1.61	2.98	0.91	0.81	88.0	7 6	ò.	0.5	98.0	1.16	1.33	0.42	9 00	9.0	C.8.	0.62	1.17	1.03	0.47	1.18	0.67	0.24	0 80	88	3	2.00	, 6	10.18	4.0	6.0	99.0	4 5	1.12	70.0	77.	6.60	4 4 5	1.28	5	1.25	85.0	10.4	5	 A	) 30 30	6.9	1.14	و.00		
		3.63	0.42	3 03	9 6	5.03	0.13	0.72	0.88	990	127	4 5	6.6	26.0	0.70	4.46	0.77	0.59	3.69	4.80	3.28	. C		77.0	200	3.31	1.14	50.	3.89	0.33	1.02	3.38	44.	3.53	4.15	0.71	3.26	3.38	3.75	7.12	3.75 9.00	3.32	i (i	3.14	3.29	3.21	3.59	3.34	0.64		
		1.10	9	B 10	0.87	- - 34	0.76	0.93	1 29		3.5	1.25	0.61	1.38	0.90	0.80	05.0	22	740		27.	7.0	1.13	99.0	3.07	0.94	0.92	1.02	0.81	0.52	0.83	0.84	1.21	0.71	0.56	0.53	1.21	1.01	<b>1</b> .	3.00	1.01	<b>79</b> .0	1.63	0.79	0.65	0.91	0.90	0.85	0.92	,	
		74.0	4.0	0.15	1.55	1.18	00 6	3 6	3.40	0.00	3.63	09.0	3.19	1.37	9.65	3.5	5.0	2 .	41.T	0.51	0.31	0.74	1.74	5.74	1.40	66.0	3.28	0.72	0.67	3.32	20.4	07.0	1.34	49.5	0.67	1.45	1 28	10.1	0.79	0.23	0.86	0.56	3.77	0.79	0.59	0.51	0.74	45.0	1 03	5 4	1/
			4.	44	1 44	ţ ;	<u>.</u> ;	4	<del>-</del> <del>4</del>	<u>-</u> .	1.44	1.43	5 F	<u> </u>	<del>2</del> .	1.43	1.43	1.43	1.43	1.42	1.42	1.42	1.42		4 5	4 6	24. 24.	4.	24.1	7 5	1.42	1,42	4.	<b>9</b> .	. ·		4.	7		17.			07	7	) · ·	9.	Q+.	1.40	04.	1.40	
			141	; ;	07.77	50.F	156.38	43.22	0.48	0.22	2	2 5	1.87	0.26	7.16	1.03	9.44	0.83	0.49	29.18	14.7		20.0	87	9.12	7.83	120	0.78	2.26	10.38	25.35	0.39	2.54	0.92	3.02	2.74	4.75	1.40	2.44	70.1	18.21	2.76	41.47	2.13	5.29	10.48	0.14	3.04	33.92	99.0	
			5	77.	<u>2</u>	1.58	25.26	62.25	69.0		2.3	<u>.</u>	2.68	0.37	10.24	1.47	190	¥	7	. 42	3 6	9 9	5 5	10.37	12.96	11.19	1.70	1.19	3,20	14.70	35.90	0.55	3.59	<b>1</b> .30	4.25	3.86	6.70	1.97	3.4	_ 8	25.61	3.88	58.28	2.99	7.43	14.69	0.20	4.21	47.45	0.92	
-7 SINE I				N67/00	AA872383	AA455659	AA683050	0707A	9163386	000000	W80067	T97890	R76505	AA099820	190980	AA974971	DIGUER	44477244	120047	1007	AA/00240	AA625890	AA677457	R25823	N47316	H87106	N75318	H99215	R43485	AA704332	AA668470	R42533	AA707121	AA448599	AA001199	AA284634	173558	AA480906	N51740	H60696	T62040	N58052	R07196	AA432124	AA706964	N24155	AA481052	· H62529	H92216	AA460282	
				291426	1472735	814018	014267	200	25	290162	416081	121530	143759	489800	443483	70571	200001	24815	757451	239446	392607	744944	460218	36393	280528	252382	298779	261851	32495	383706	957878 908578	30275	452059	785975	162251	713193	82734	814595	281737	208985	85609	247446	126847	781467	451918	269563	814661	208165	221778	795730	,
				713	717	, ,	2 1	2	717	718	719	2	i i	į	3 1	3	124	725	728	727	728	729	2	Ē	É	1 5	3 5	Ş	3 5	3 6	ž ŝ	2 1	P (	₹;	₹ ;	7 5	2 ;	3,5	7.48	777	7.48	2 %	7.65	3 5	<u> </u>	, F	3 2	? ¥	3 2	3 5	č

Table 2-7: Good vs. Poor Clinical Outcome: Genes upregulated 3 fold or more in at least 1 tumor sample.

<del>-</del> -	-	-	<b></b> .	-	<b>-</b> -	<del>-</del>	-		-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	_		- •	-		
0.81	1.1	0.78	1.05	1.01	1.12	1.08	0.75	99.0	1.03	4.05	1.21	0.38	1.10	0.48	9.0	0.78	1.58	1.10	3.68	0.72	0.68	0.71	0.87	1.28	1,90	0.70	0.73	1.00	0.83	0.98	0.64	1.28	4.1	0.86	3.18	1.40	1.18	0.43	0.47	60	98.0	5 6	<b>.</b>			
0.71	8.	0.80	0.80	0.97	1.08	0.63	0.61	1.29	0.79	0.50	0.78	0.39	1.25	3.57	77.0	0.68	790	4 30	÷ 5	10.0	200	20.5	2	94	0.65	0.88	1.77	0.38	0.70	4.06	1 15	0.93	1.24	0.47	0.89	0.31	1.39	0.30	25.0	, c	600	0 i	1.73	0.99		
3.16	3.62	3.10	3.47	3.60	3.47	3.24	0.52	3.07	3.29	69.0	3.20	<b>4</b> .28	0.24	22	89	8 6	8 6	3 5	÷ 6	- 8:0 • 4:0	.03	2.85	90.0	3.20	9 6	9.00	7 O.02	5.08		c	9 6		30.5	78.0	530	98.6	20.E	5 7	5 2	e (	91.5 7.1	0.91	3.16	3.07		
1.56	72.0	* 0°.	5 C	0.75		6.0	. ÷	90.0	9.0	, e		5.5	0.40	0.73 8.73	3 6	5 c	0.0	26.0	c/.n	F. 6	3.06	25.5	20 E	0.79	0.82	98.0	80.0	S :	3.17	3.3/	0 0 4 }	2.5	- P	7.0	4 4	20.0	/6.0 7.0	<b>\$</b> 60.0	98.0	F.7	0.86	0.95	0.77	0.83		
0.75	9 6		0.0	0.73	3 6	) C	, c	9 9	0.80	) S.O.	94.0	0.62		- CT	J. 1	0.61	3.81	0.66	0.57	0.85	0.36	0.48	0.80	0.83	0.72	3.05	3.53	0.26	0.50	0.77	9.0	1.20	0.50	Q. (	0.0	11.1	0.08	0.45	0.59	0.43	0.71	3.41	0.49	0.78	18	)
1.40	1.40	1.39	1.39	1.39	1.39	1.39	1.39	1.39	1.38	1.38	1.38	1.38	1.38	1.38	1.38	1.38	1.38	1.37	1.37	1.37	1.37	1.37	1.37	1.37	1.37	1.36	1.36	1.36	1.38	1.36	<del>.</del> 36	1.36 	1.35	1.35	1.35	1.35	1.35	1.35	1.34	1.34	<del>.</del> .	75.	7	<u> </u>	<u> </u>	
6.17	15.21	19.57	0.00	1.32	¥ ¥	52.40	1.03	0.42	1.01	2.08	2.42	0.85	1.27	1.29	2.02	0.73	0.52	0.57	102.66	0.10	13.04	1.68	0.70	3.31	13.58	1.57	0.92	7.82	26.33	2.60	1.37	2.32	19.13	24.47	<b>9</b>	1.61	0.13	51.24	0.67	1.16	5.76	49	99.0	6.0	5.5	
8.62	21.24	27.27	0.13	<del>2</del> .	48.51	72.84	1.43	0.59	1.40	2.88	3.34	1.17	1.78	1.79	2.79	1.01	0.71	0.78	140.80	0.13	17.85	2.29	96.0	4.52	18.54	2.14	1.25	10.64	35.81	3.53	1.86	3.15	25.91	33.12	0.08	2.18	0.17	69.06	0.91	1.56	7.73	2	200	2.0	D 4.0	
R78607	H84759	AA702193	AA054439	AA598775	H81716	H65300	H04757	R17747	AA678361	R56840	R00707	AA682565	N92895	R44837	AA430367	AA430410	R19878	AA704749	H94870	AA776702	N62394	AA682642	AA431772	R70598	AA279172	AA620608	AA443116	AA291972	AA425692	AA447764	AA434403	A428341	H92215	M454825	AA233549	N36402	H86275	AA708348	AA857496	R45118	H79078	AA253442	HOTETA	0.07747	028444	
144932	219937	384134	381054	898058	219861	208225	152270	25396	431866	138255	123474	431269	309081	33854	769857	769947	34888	451161	230236	1292829	28863	450580	782501	141854	704076	1048698	809488	725364	773204	813661	770879	769673	221778	814744	666128	273048	223314	392673	1384851	35010	248724	£40124	00830	27/152	32285/	
158	759	92	761	762	763	764	765	88	787	<b>168</b>	769	2	<u>=</u>	E	E	14	77.5	1,18	E	*	2 6	780	ž.	5 2	. £	¥ 5	785	788	787	788	789	280	791	792	8	ğ	795	28	797	8	2 6	8	3 3	<b>S</b>	802	

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0.33	1.85	1.05	3.01	1.14	0.65	3.11	0.70	0.86	0.59	0.65	3.43	9.0 89.0	270	3.4	- c	30.0	200	\ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \	5 . c	5.5	1.15	1.5	50.1	080	0.82	0.98	09.0	0.87	1.16	0.75	1.20	0.86	3. 3.	) (C. 0.	07.	C . C	70 C	0.02	8 8	3 5	2.0	86.0		
0.58 1.47	0.62	0.82	1.00	1.01	0.51	1.02	3.43	4.10	0.55	0.82	0.77	1.13	0.95	1.05	76.0	0.43	21.1	0.85	0.76	1.33	2 2	1.0	(S, C)	9 C	7.07		- 25	1.17	1.02	0.68	0.48	0.68	0.57	0.91	0.78	0.88	0.41	1.03	0.76	0.87	4.0	0.38		
4.83 3.01	6.34	3.50	0,67	3.45	3.26	98.0	0.97	0.45	0.51	3.86	0.46	3.04	3.63	3.40	0.50	0.5%	3.16	3.49	0.23	3.39	0.89	3.01	0.31	3.25	0.62	5. 6. 6. 6. 6. 6. 6. 6. 6. 6. 6. 6. 6. 6.	0.67	) (1) (2)	5 6	3.29	021	3.46	3.77	3.32	0.57	3.06	3.87	3.30	3.32	3.06	3.30	0.93		
0.57 0.57	3 32	1000	6	. 6	6.5	7.5	89	200	25. 25.	S 6	200	67.0	45.0	0.70	4.0	4.33	9.0 29.0	0.70	0.73	0.56	0.81	69.0	0.93	0.67	9.0 <b>2</b> 9.0	0.80	4. 4.	0.61	0.62 0.63	9 6	9 6	0.86	1.08	0.74	0.66	0.62	0.77	0.75	0.58	0.60	8	3.67		
0.35		0.52	9.0	2. 6	14.0	D 0		8.5	4.0	0.4	0.42		0.7	55.0		0.25	0.61	0.63	3.68	0.68	3.04	0.63	3.14	45.0	3.76	0.78	3.02	0.63	0.74	0.60	0.99	3.10	2.00	5 6	3.13	2 5	. 55.0 55.0	95.0	92.0	88.0	0.00	55.0	<b>1</b>	<u>D</u>
1.33	1.33	1.33	<del>ا</del> .3	1.33	1.32	1.32	1.32	1.32	1.31	1.31	1.31	<del>.</del> .3	1.3	1.5	5.5	5.	5	5.	1.30	5. t	5. 5	5 5	5.5	25.	22	2	1.28	1.28	1.28	1.28	1.28	1.27	1.27	1.27	1.27	1.27	77.	07.1	07:1	07.1	1.20	97.1	67:1	
2.54	78.95	1.93	26.74	0.52	4.55	0.10	0.38	0.63	2.29	11.10	0.64	1.12	0.50	5.91	11.97	<b>*</b> 5	11.48	3.82	10.08	85.4 80.4	21.1	17.7	37.78	6.50		. e	8 6	1.15	3.40	35.99	9.65	2.08	0.21	2.88	0.32	9.82	23.43	2.13	1.28	2.16	17.73	S 1	1.72	
3.38	105.05	2.57	35.55	0.70	8.02	0.13	0.51	0.83	3.01	14.58	8.0	1.46	0.65	<b>5.</b>	15.65	5	14.99	4.99	13.13	5.71	7.45	2.88	48.29	2.66	12.02	?	- 5	? ?	4.47	46.13	83.0	5.66	0.28	3.67	0.40	12.44	29.64	2.68	<b>.</b> 8	2.72	22.29	7.0	2.15	
AA676920	W37532	AA497111	H66875	H09172	AA682838	AA758470	AA432134	AA447592	AA17921	AA873159	AA401828	AA464979	AA504505	AA976561	AA704908	AA927544	AA485303	AA677337	H68553	T63324	AA427561	R00283	N62763	H11732	AA701328	AA164830	AA/ /8063	A A A B 4 4 4 7	AA704370	AA187334	T69593	AA279060	AA485249	N47443	AA774501	AA157813	R66923	AA677300	AA482127	N50859	N76088	AA486765	N59532	
AEDIEA	272438	922130	241387	70549	450300	434020	781492	782687	767405	1477689	775274	810109	825366	1558655	461383	1552618	840266	454469	212473	80109	770069	123278	288562	47481	435126	595148	1049030	49499	130334	202/32	83541	703916	815835	280697	399264	688915	140289	454795	766450	280982	295454	641057	248631	
		<b>4</b> 6	2 2	3 5	38	3 8	8 6	2 6	2 2	4 5	2.5	2 4	816	817	818	819	820	821	82	823	824	828	826	827	828	828	<b>ස</b>	<b>3</b>	<b>3</b> 9	3 8	2 E	8 8	837	2	83	840	3	842	2	Ž	<b>2</b>	26	847	

0.76 0.96 0.60 0.88 0.76 3.03 3.12	0.68 3.01 4.02 0.00 0.73 0.73 0.73 0.82 0.78	0.74 0.63 3.05 3.05 1.34 1.04 1.04 0.20 0.20 0.32	0.23 0.52 0.00 0.05 0.72 3.09 3.09
0.40 0.61 0.05 0.88 0.89 0.27	0.00 0.00 0.00 0.54 0.54 0.00 0.00 0.00	0.000000000000000000000000000000000000	0.00 0.00 0.00 0.00 0.45 0.62 0.62
0.34 3.03 3.25 3.26 0.40 0.70 0.71	0.72 0.29 1.20 3.03 3.03 3.14 3.30 3.30	3.14 3.28 3.28 3.19 3.05 3.05 3.25 3.25 3.25 3.25	0.30 0.30 0.37 0.54 0.32 0.30 0.11
0.93 0.63 0.63 0.64 0.63 0.63 0.83	0.59 1.77 1.77 1.12 0.80 0.59 0.74 0.74 0.74	0.65 0.66 0.09 0.74 0.64 0.63 0.05 0.06 0.73	1.12 1.05 1.07 1.07 3.69 3.39 3.39 0.54 0.55 0.34
3.84 1.04 0.79 0.56 3.33 1.32 3.17	9.10 9.10 9.10 9.14 9.63 9.10 9.53 9.63 9.63 9.63	0.55 0.65 0.65 0.65 0.65 0.65 0.53 0.53 0.53 0.53	0.30 0.22 0.00 0.00 0.44 0.44 0.44 0.44
21 22 23 24 25 25 25 25 25 25 25 25 25 25 25 25 25	<b>7</b>	222 222 222 222 222 222 223 223 223 223	1.15 1.14 1.17 1.10 1.08 1.03 1.03 0.97
4.19 1.47 5.09 2.25 15.13 2.25 18.72 3.79	1,52 1,35 1,35 0,09 0,03 0,06 11,12 17,78 4,32 3,92	0.58 4.94 14.12 31.16 3.116 5.54 8.11 2.96 0.95 15.23 1.58 1.25	2.63 1.20 30.30 0.02 0.02 2.14 7.59 6.99 10.12
5.25 1.84 0.37 2.82 18.89 23.31 4.71	1.88 1.68 1.68 0.04 0.07 3.02 13.72 13.72 11.57 4.80	0.71 6.03 17.18 37.89 6.64 9.69 3.53 7.79 1.10 17.56 4.23 15.02	3.01 1.36 34.68 0.03 0.02 2.31 7.22 10.45
R34225 A4491251 A4707080 A4213667 H86559 A4451863 A4451863	AA465194 H54384 AA001444 N32071 AA426288 AA426288 AA701978 AA701978 AA701978 AA701978 AA701978 AA701978	AA703159 H38848 AA504132 N21688 A1015196 R34273 AA704278 AA4699719 R16146 AA466495 R01304 H10072	AA628979 R64454 T63245 AA057195 AA131885 AA521384 N49107 AA625668 AA478470 N91767
136026 824270 451596 683274 223180 786308 196189	815090 203350 361943 260168 1404985 773284 435953 480143 52730 270331 40636 148838	435596 190692 82524 268455 1623328 136070 450997 433300 53158 814053 123755 48843 754280	745400 141361 80050 489076 504358 826995 279800 745347 786602
848 849 850 851 853 853	858 857 853 860 861 864 865 865 865 865	869 870 871 872 873 874 875 876 876 877 878	882 884 884 885 885 887 887 890 890

Table 2-8: Good vs. Poor Clinical Outcome. Genes upregulated 3 fold or more in at least 2 tumor samples.

Table 2-8: Good vs Poor Clinical outcome: Genes upregulated 3 fold or more in at least 2 tumor samples

50 - 100 4 10 10 1 20 1 20 1 20 1 20 1 20 1	1				Folial Average OCCAV		olo(mpinser.			FORTANGOODIAY FOLD MININGSTAN	Saldmes
- 28 4 58 6 7 8 6 6 7 7 7 7 7 7 7 7 8 6 7 7 8 6 7 7 7 7		Accession	AveGood	AvePoor	ePoor)	ePoor)	ePoor)	ePoor)	eroor)		
- 1		Number				r	08.9	3.51	99'0	7.30	4
. スキャゥト 890000000000	299679	W05628	2.55	0.50	5.07	70.7	9 5	6.33	4.17	6.26	4
	811162	AA486471	28.87	6.85	£.24	D.'O	5 6	4 22	2,10	21.15	es .
**************************************	171641	AA034945	11.03	1.08	10.17	21.19	7.70	76.0	9.85	7.08	က
	281039	N47717	36.67	4.42	8.29	22.87	7.40 4.40	73.0	7.05	8.80	က
0 8 7 8 8 9 7 7 7 7 7 7 8 8 9 7 8 8 9 7 8 8 9 7 7 8 8 9 7 8 9 9 7 8 9 9 7 8 9 9 9 9	282089	N48259	27.13	3.45	7.87	22.41	- C	9 5	9.52	15.90	ຕ
0	155072	R71393	3.32	0.45	7.44	8.50	9.5	9 8	530	10.17	က
× 8 9 5 11 12 12 14 15 15 15 16 16 16 16 16 16 16 16 16 16 16 16 16	208210	H65286	2.70	0.42	6.48	14.00	08.1 1.00	8.0	930	5.96	ю
0 8 9 7 7 7 7 7 7 7 8 8 8 8 8 8 8 8 8 8 8	770068	AA430545	26.72	4.62	5.78	12.05	/9:0 7 0 7	0.5	4.37	12.92	9
0 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	712950	AA282273	38.22	<b>8</b> .9	5.50	8.84	2 5	5.0	85	3.28	က
22224544545	81331	T60111	4.13	0.82	5.03	11.18	0.50	979	3.86	5,42	m
22545545555	287728	N59158	3.49	0.71	<b>6</b> .90	14.39	0.35	4.46	2.14	3.71	က
1275272525	41228	R58953	0.05	0.01	4.38	9.07	7.33	0,4	000	3.18	m
7 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5	122345	199191	0.03	0.01	4.16	12.66	7.7	133	8,3	9.31	က
15 16 18 19 19 19 19 19 19 19 19 19 19 19 19 19	186918	H43317	6.27	1.60	3.83	3.28	14.6	7.7	2.55	5.03	eo .
25 25 25 25 25 25 25 25 25 25 25 25 25 2	1049033	AA778575	2.50	0.65	3.87	98.4	7.70	88	3.51	6.23	ო
<b>1289</b>	195365	R88904	1.81	0.50	3.62	6.03	0 7	7.0	2.28	0. 2.	က
2522	50114	H16743	1.48	0.43	3.43	5.50	÷ ÷	122	1.3	1.35	ო
2823	638611	AA456975	87.41	25.99	3.36	L'6	7 6	1.42	3.24	3.77	က
ឧក	283196	N45282	5.16	1.55	3.32	1.57	5.0	7	4.76	3.74	က
<b>7</b> 7	745525	AA626248	4.16	1.27	3.28	0 0 0 0	1.7.	07.6	1.59	3.11	ന
; ;	159487	H15926	0.12	0.0 20.0	2.82	2.50	, c	2.87	0.79	3.97	က
?	103473	AA779728	76.12	27.06	2.81	2.22	3.41	56.	3,32	3.28	ന
ឧ	13844		2.02	0.74	2.75	4.16 0.00	8 6	- C	3,13	3.31	ന
7	378488	<b>A777187</b>	17.01	7.17	2.37	3.23	1.40	28.44	49	1.02	7
8	767130	<b>AA2</b> 4535	4.28	0.37	11.65	09.1	74.7	44.45	39.	1.22	7
78	307337	N95226	10.81	0.96	11.29	60.1	4 4	33.58	2	3.17	2
27	502625	AA134576	7.27	0.9	8.08	1.44	 	15.74	S:	0.71	6
78	41889	R86438	1.41	0.28	50.c	D 10	2 7.5 7.7.5	2.83	0.99	7.09	7
82	301735	N90882	4.10	0.80	4.58	20.6	3.53	11.13	<del>2</del> .	2.65	М
ន	136605	R35051	32.22	7.39	4.36	6.73	3	0.78	6.56	7.23	7
ភ	382278	AA001219	5.53	¥.	3.61	\$ 63 63 63 63	4 50	10.09	2.89	1.01	7
33	756708	<b>AA443903</b>	2.58	0.87	10 m	0.30	. 4 . 5	11.96	0.59	0.98	7
ន	265028	N21084	1.23	0.32	3.78	90.0	, r,	11.75	0.59	0.31	74
ಸ	1409509	_	284	0.75	3.78	0.30	9 6	10.11	1.36	3.00	8
35	365707	AA025434	1. 18.	0. S3	3.68	6.0	S 4	10.03	0.93	0.97	N
8	768953	AA424920	3.43	0.94	3.66	9 (9	. v	0.68	2.00	9.41	7
37	357628	W94121	1.80	0.49	3.65	2.62	 	2 07	¥.	1.38	7
೫	35058	R45192	28.75	7.80	3.64	n n	į				

Table 2-8: Good vs. Poor Clinical Outcome. Genes upregulated 3 fold or more in at least 2 tumor samples.

ппппппп	0000000	,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	, , , , , , , , , , , , , , , , , , , ,	папапапапапа
9.44 6.17 4.57 2.75 0.88 3.90	8. 4. 9. 9. 9. 9. 9. 9. 9. 9. 9. 9. 9. 9. 9.	0.09 5.4.4.5 6.4.7.4.4.4.4.4.4.4.4.4.4.4.4.4.4.4.4.4.	4.89 9.44 9.34 1.24 0.72 4.89 9.42 1.06	1,03 3,11 6,51 0,63 3,26 1,19 1,112 3,25 6,89 3,86
5.51 8.90 6.90 0.99 2.25 4.65	2.50 2.42 2.42 2.43 0.68 0.49	3.20 0.60 1.70 1.70 1.58 0.82	5.48 2.55 0.46 0.69 0.69 1.56 2.10 4.00	3.15 3.69 5.65 1.21 1.76 1.58 1.13 2.28
0.73 1.32 2.38 5.00 10.28	2.26 1.26 0.59 0.45 1.80	0.57 0.74 4.46 0.94 1.52 3.16 6.22	0.79 1.14 1.55 7.31 7.85 1.62 1.71 0.54 6.17	5.53 2.58 1.09 5.67 2.12 1.95 1.56 0.22 0.22
0.52 2.44 1.91 2.54 3.45	2.96 2.96 1.26 0.81 0.71	0.89 4.10 2.99 1.13 1.41 6.21 3.75 3.46	1.98 0.96 1.25 1.25 0.54 1.22 1.22 1.46 0.37	2.15 0.93 0.93 3.19 3.72 3.10 0.53 3.38 3.78 1.42
1.47 1.63 1.01 3.72 0.86 8.38	0.59 6.07 8.02 7.62 3.86 6.21	0.04 0.74 0.74 0.82 0.92 1.83	4.84 0.63 0.63 3.32 3.32 3.32 3.42 1.63 3.44 4.42	2.54 2.17 2.17 0.44 1.63 0.61 3.24 0.31 0.80 0.70 3.36 2.37
8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8	8288 444 448	3.09 3.09 2.28 2.28 2.27 2.77 3.75	2 2 2 2 2 2 2 3 3 3 5 6 6 6 6 7 3 3 5 6 6 6 6 7 3 3 5 6 6 6 6 6 6 7 3 3 5 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6	2
8.55 0.02 1.14 0.79 3.68	2.48 0.25 1.18 1.88 0.98	0.02 0.75 1.36 0.64 36.00 0.75 35.15 0.45	17.94 17.94 17.94 1.09 1.00 1.10 1.17 2.46	0.94 1.63 15.73 15.73 1.38 0.48 0.57 1.52 1.52 1.53
30.23 0.08 13.79 3.83 2.60 12.10	8.05 3.79 7.05 5.91 4.04	0.06 2.32 4.17 1.82 101.29 2.12 98.80 1.23	46.24 39.72 3.37 2.93 6.96 1.50 2.62 3.92 6.38	2.44 4.21 0.61 39.31 1.87 3.37 1.16 52.42 1.37 3.93 3.93
R77948 AA43867 AA633658 AA412417 R66101 H09076	R00276 R61372 AA628113 N5347 AA775257 T87128 AA32270	H47076 H02837 AA418748 R45517 N55459 N59816 AA199979 AA700880	A44592 7446792 76281 76947 76947 A449698 N53031 A4150500 76305 W7433	H17121 AA702335 R61883 H72722 AA453526 AA453526 AA534031 H23340 H23340 H23590 H73590 AA77955 H73590
145740 757200 856575 731469 140197	123264 37980 1055707 245277 878596 66599 782335	193087 151477 768020 35329 245990 288983 646037 434952	2315.4 840590 78353 1418621 66902 823655 246430 491751 82905 346583	51178 447416 42872 232772 277173 785425 743868 743868 74000 202535 267546 740604 214441 364022
862444	54 4 4 8 6 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5	28 27 28 28 28 28 28 28 28 28 28 28 28 28 28	5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5	7 7 7 7 7 7 7 7 8 7 8 8 8 8 8 8 8 8 8 8

Table 2-8: Good vs. Poor Clinical Outcome: Genes upregulated 3 fold or more in at least 2 tumor samples.

77	7	7	7	7	2	7	7	2	2	2	7	7	2	74	2	7	7	2	8	~	7	2	7	7	7	7	7 (	71	7	7 (	۷ (	<b>4</b> C	4 6	۰,	4 (	1 C	4 6	4 6	<b>v</b> (	NE (	N 6	<b>J</b>		
1.24	1.96	4.34	3.45	0.0	1.87	4.40	3.90	1.32	5.80	4.05	4.08	4.96	3.89	1.53	3.95	0.92	3.62	3.44	4.38	3.80	1.03	3.61	3.34	3.05	4.26	4.08	3.30	3.07	 	3.16	3.85	4.12	4.24		3.05	0.0	9.79	3.13	= :	0.61	0.49	0.48		
3.44 2.96	3.80	0.18	4.10	0.74	4.69	0.24	4.70	3.63	0.51	5.78	50.50	0.23	4.06	0.71	6.39	1.60	1 67	2.03	3.46	1.07	90	4.0	28.0	2.36	0.40	4.51	3.19	3.09	0.93	3.41	3.03	0.54	0.58	3.49	0.41	0.75	3.65	3.37	3.19	0.56	0.63	<b>7</b> .04		
1.42	40.4	S 6	8 6	33.0	6,0	1 1	6	5 C	. c	5 6	2 9	0.35 75	£ 5	250	9,0	0 K		9.00		7 2	5 5	3.02	5 6	5.5	1.13	0.52	0.98	1.23	0.68	0.93	0.78	0.37	0.38	0.85	1.87	4.05	0.52	0.21	3.72	3.89	3.81	3.25		
1.57	F	5.55 6.40	2.10	0.00	- c	5 G	0.00	5.6	20.0	9.78		0.52	1.4.0	) (S		94.0 20.0	50.0	1.35 5.5	0.50	0.67	C.9.	1.92	5.00	6.33	3.23	) E	69	105	60	0.93	0.76	28.5	3.02	0:30	3.19	3.33	99.0	1.44	0.56	3.19	3.18	0.53		
3,94	3.32	3.12	4.65	2.81	4.73	3.34	9/0	0.56	3.06	3.40	0.17	0.32	1.03	3.28	4.01	0.34	1.59	0.89	3.61	0.76	3.99	3.33	0.84	3.08	90.0	0.20	0.20 0.76	5.5		2 5	3 4	260	52	0.74	0.91	55	0.42	690	2,56	750	98.0	00.0		ო
2.32	2.31	2.30	2.27	2.26	2.23	2.23	2.22	2.22	2.20	2.19	2.19	2.18	2.17	2.17	2.14	2.13	2.11	2.11	2.10	2.10	2.09	2.07	2.06	2.08	2.04	2.01	3. G		9 2	) S	8.0	8 8	68	8	1.89	8	1.80	1	14.	7.	0/:	. 99	<b>?</b>	
0.33	<del>.</del> 2	0.37	<b>6</b> .85	1.26	0.70	2.14	12.67	5.54	1.40	17.15	15.57	13.39	2.98	0.24	1.57	10.50	1.7	3.65	22.41	0.86	0.30	0.83 28.0	10.94	9.27	3.81	0.0	25.44	2.07	1.1.	1.50	C.83		2. 18 2. 2.	87.0	25.52	13.84	124	24 24	27.24	£ . 73	\$ 75°		, ,	
0.77	2.41	0.84	15.54	2.85	1.57	4.78	28.11	12.27	3.07	37.52	34.05	29.18	6.47	0.52	3.35	22.38	3.60	7.69	47.09	1.81	0.62	Ē.	22.57	19.09	7.78	0.02	50.71	4.12	2.20	2.08	1.67	7.5	, , ,	3	36 95	26.33	£3.14	3.5	07.70	46.33	2 9		3	
N67039	AA447522	H08862	T67053	N52151	AA456299	AA456321	AA663981	AA988345	R76247	H16803	H77766	N57872	AA884926	H50086	AA434487	R16596	W49563	N72137	AA070226	AA779457	H15677	AA809586	AA111969	AA668897	AA683520	AA459292	N80129	AA035558	R36989	AA459197	AA040424	N64/06	NAZONO	A A 457528	W7024	Naces	A 4 80666	A A 630006	200000	1000	A418000	K46000	A10000A	
295985	782575		66560	284383	813151	_		_	144747	50772	214162	247117	_			129585	324815	291097	530814	-	49303	1031701	530185	863398	378813	810899	297392	471682	27098	810873	485854	283292	28833/	00340	34300	79780	238300	824340	884606	132702	767449	35804	20113	
2	55	88	87	88	89	8	2	85	83	ā	98	8	97	86	68	8	101	102	50	\$	105	\$	107	108	109	5	Ξ	12	113	7	15	<del>1</del>	11	0	118	07.	5 5	7	123	124	52	128	171	

Table 2-9; Good vs. Poor Clinical Outcome: Genes upregulated 5 fold or more in at least 2 tumor samples.

Ordari         HAADE Close ID HAADE HARDEN         One Bank HATTI HAADE HARDEN         Tissue HARDEN         Tissue HARDEN         Tissue HARDEN         Avedon HARDEN         Folicityword Avedon HARDE	ı		tumor samples.	mples.									
281030         HAF717         Done         38.87         4.42         8.28         7.87         22.41         0.41         0.27           282020         HAR529         proxisiae         27.13         3.45         7.87         22.41         0.41         0.67           155072         HAR529         proxisiae         27.13         3.45         7.44         8.50         1.97         1.30           208210         HAR528         brain         2.02         0.45         7.44         8.50         1.97         1.30           208210         HAR528         brain         2.02         0.45         5.78         1.40         0.69           770066         AAA30446         brain         1.03         0.42         5.78         1.205         0.87         1.50           770066         AAA30446         brain         1.03         0.46         5.77         0.80         2.84           770066         AAA24536         kidniny         4.28         0.37         11.65         1.80         2.84         4.45           771240         AAA24636         brain         1.031         0.89         1.129         1.80         2.84         1.44           771441         <	- Pro	IMAGE Clone (D	Gen Bank Accession	Tlasue	AveGood	AvePoor	Fold(AveGood/A	. Fold(mpm27/A vePoor)	Fold(mpm284A vePoor)	Fold(mpm29/Av ePoor)	Fold(mpm30/Av •Poor)	Fold(mpm3&/Av •Poor)	# OF Tumor Samples
202009         NA6259         prostate         27.13         3.45         7.87         22.41         0.41         0.67           155022         PMS259         prostate         27.13         3.45         7.44         8.50         1.97         1.30           15502         H65266         testis         2.70         0.42         6.46         14.00         1.98         0.98           770066         AAA30646         brain         2.67         6.76         5.77         0.87         1.90           290670         W058226         brain         1.68         0.37         11.65         0.87         1.90           290679         W058226         brain         1.0.61         0.86         5.77         0.89         3.51           787130         AAA24636         kidney         4.28         0.37         11.65         1.60         3.51         1.90         0.89         28.44         1.90         0.89         28.44         4.44         4.44         4.44         4.44         4.44         4.44         4.44         4.44         4.44         4.44         4.44         4.44         4.44         4.44         4.44         4.44         4.44         4.44         4.44         4.4	-	281030	NATA 1		38.87	4.42	8.29	22.87	1.40	0.27	9.85	7.08	
155072         R71303         3.32         0.45         7.44         8.50         1.97         1.30           203210         H62266         testils         2.70         0.42         6.48         14.00         1.98         0.98           770066         AAA30646         brash         26.72         4.62         5.76         12.05         0.67         1.90           298679         W05628         borne         2.56         0.60         5.07         7.07         6.80         3.51           298679         W05628         brash         11.03         11.65         11.60         25.69         28.4           307337         N89226         brash         11.03         10.61         11.29         1.65         7.47         44.45           30734         N89226         brash         11.03         10.61         11.29         1.66         7.47         44.45           471849         Brash         brash         1.41         0.28         5.03         1.44         5.00         4.22           471849         R64340         brash         1.41         0.28         5.03         1.44         5.00         1.44         5.00         4.22           41869 <td>. ~</td> <td>282089</td> <td>N48259</td> <th>orostate</th> <td>27.13</td> <td>3.45</td> <td>7.87</td> <td>22.41</td> <td>0.41</td> <td>0.67</td> <td>7.05</td> <td>8.80</td> <td>က</td>	. ~	282089	N48259	orostate	27.13	3.45	7.87	22.41	0.41	0.67	7.05	8.80	က
20810         H65266         Lestis         Lestis         2.70         0.42         6.46         14.00         1.98         0.88           770066         AAA30646         brain         29.72         4.62         5.78         12.05         0.67         1.90         0.88           298670         WOSG2A         borne         2.55         0.50         5.77         7.07         6.80         3.51           787130         AAA2463         kidney         4.28         0.37         11.65         1.60         25.69         28.44           307337         N83226         brain         11.03         1.08         11.29         1.85         7.47         44.45           471841         AAA2453         kidney         4.28         0.37         1.185         1.47         44.45           471840         AAA22273         t-ceil         36.22         6.64         5.50         8.84         1.04         0.35           41869         Re643b         actennal         1.41         0.28         5.63         1.46         5.93         1.47         44.45           287728         AA32420         actennal         1.41         0.28         5.63         1.46         5.93	m	155072	R71393	,	3.32	0.45	7.4	8.50	1.97	1.30	9.62	15.90	e
770066         AAA30646         brash         28,72         462         5.78         12.05         0.67         1.90           289670         W05625         bore         2.55         0.50         5.07         7.07         6.80         3.51           787130         AAA24635         kidney         4.28         0.37         11.65         1.60         25.69         28.44           30737         N8226         spleen         10.61         0.86         11.29         1.65         7.47         44.45           30737         N82227         spleen         10.61         0.86         10.77         21.19         22.0         28.44           471869         R6443         5.60         8.84         1.04         0.35           41869         R6443         5.60         8.84         1.04         0.35           41869         R6443         5.60         8.84         1.04         0.35           41869         R6443         5.60         8.84         1.04         0.35           287728         M59158         4.13         0.82         5.03         1.46         5.80         1.57           287728         M59188         4.10         0.80	•	208210	H65286	tests	2.70	0.42	6.48	14.00	1.98	86.0	5.30	10.17	60
299679         WOSGLS6         borne         2.55         0.450         6.07         7.07         6.80         3.51           787130         AAA24635         kidney         4.28         0.37         11.65         1.60         25.69         28.44           307337         N89226         spleen         10.61         0.86         11.29         1.65         7.47         44.45           471240         AA034046         brahn         11.03         1.06         10.17         21.19         2.20         4.22           471240         AA034046         brahn         11.03         1.06         10.17         21.19         2.20         4.22           471260         AA282273         t-cell         36.22         6.84         5.50         8.84         1.04         0.35           41869         R66436         adrenal         1.41         0.28         5.03         11.46         5.89         15.74           41869         R66436         AA282273         AA486471         A4467         0.28         4.19         0.80         4.46         0.30         1.57         4.45           28772         AA486471         Infryride         28.87         4.26         0.76         2.	vo	770066	AA430646	prain	26.72	4.62	5.78	12.05	0.67	1.90	8.30	5.96	e,
787130         AAA24635         kixliney         4.28         0.37         11.05         1.66         25.69         28.44           307337         N89226         spleen         10.61         0.66         11.29         1.65         7.47         44.45           471641         AAO34046         brah         11.03         1.08         10.17         21.19         2.20         4.22           471640         AAO34046         brah         11.03         1.08         10.17         21.19         2.20         4.22           41869         R66436         adrenal         14.41         0.28         5.03         1.46         5.93         15.74           41869         R66436         adrenal         1.41         0.28         5.03         1.46         5.93         15.74           287728         AS6486         1.43         0.82         5.03         14.36         0.36         0.46         0.80         1.57         1.80         0.46         0.80         1.80         0.46         0.80         1.80         0.46         0.80         1.74         0.76         0.80         1.74         0.76         0.80         0.46         0.80         0.46         0.80         0.46         0	•	299679	W05628	pore	2.55	0.50	5.07	7.07	6.80	3.51	99.0	7.30	m
307337         N89226         spleen         10.61         0.86         11.29         1.65         7.47         44.45           471641         AAQ3446         brain         11.03         1.08         10.17         21.19         2.20         4.22           712950         AAZ82273         t-celf         36.22         6.84         5.50         8.84         1.04         0.35           41890         R66430         adrenal         1.41         0.28         5.03         1.46         5.90         15.74           28772         R66430         adrenal         1.41         0.28         5.03         1.46         5.93         15.74           28772         R66430         adrenal         1.41         0.28         5.03         1.43         0.36         0.36         0.46           28772         AAA88647         thyrold         2.84         0.75         3.61         0.70         3.61         0.75         2.83           36272         AAA001219         lymph         3.53         1.41         3.81         3.64         0.36         5.41         0.78           140850         AAA24920         Lona         2.84         0.75         3.65         0.36 <th< td=""><td>1</td><td>787130</td><td>AA424635</td><th>kidney</th><td>4.28</td><td>0.37</td><td>11.65</td><td>1.60</td><td>25.69</td><td>28.44</td><td>1.40</td><td>1.02</td><td>7</td></th<>	1	787130	AA424635	kidney	4.28	0.37	11.65	1.60	25.69	28.44	1.40	1.02	7
471641         AA034046         brahn         11.03         1.08         10.17         21.19         2.20         4.22           712950         AA202273         L-cell         36.22         6.94         5.50         8.84         1.04         0.35           41869         R86436         adrenal         1.41         0.28         5.03         1.46         5.93         15.74           81331         T60111         4.13         0.82         5.03         11.18         1.30         0.80           28728         Ms9158         3.49         0.71         4.90         14.39         0.35         0.46           301735         Ms0682         esophagus         4.10         0.80         4.56         9.05         2.75         2.83           811162         AA486471         thyrold         28.87         6.85         4.21         0.70         3.61         6.33           382278         AAA001219         lymph         5.53         1.41         3.91         2.64         2.34         0.78           1408509         AAA24220         borne         2.84         3.66         0.36         5.34         0.78           18636         AA432270         prostate	•	307337	N85226	spleen	10.81	0.08	11.29	1.65	7.47	44.45	39.1	12	7
712950         AAA32273         1-cell         36.22         6.84         5.50         8.84         1.04         0.35           41869         R66436         adrenal         1.41         0.28         5.03         1.46         5.93         15.74           81331         T60111         4.13         0.82         5.03         11.18         1.30         0.80           28728         M59158         3.49         0.71         4.90         14.39         0.35         0.46           301735         N90682         esophagus         4.10         0.80         4.56         9.05         2.75         2.83           811162         AAA48471         thyrold         28.87         6.85         4.21         0.70         3.61         6.33           382278         AAA001219         hymph         5.53         1.41         3.91         2.64         2.34         0.78           1408509         AAA24020         AAA24020         3.43         0.84         3.65         0.36         5.41         10.03           35763         R88604         1.81         0.60         3.65         6.03         1.45         0.88           145740         R77946         protate         <	œ	471641	AA034946	brah	11.03	1.08	10.17	21.19	2.20	4.22	2.10	21.15	7
41869         R66436         adrenal         1.41         0.28         5.03         1.46         5.93         15.74           28772         M58156         4.13         0.62         5.03         11.18         1.30         0.80           28772         Assesse         esophagus         4.10         0.80         4.56         9.05         2.75         2.83           301735         M90852         esophagus         4.10         0.80         4.56         9.05         2.75         2.83           811162         AA486471         thyrold         28.87         6.85         4.21         0.70         3.61         6.33           362276         AAA86829         borne         2.84         0.75         3.78         0.36         5.64         11.75           140850         AAA86829         borne         2.84         0.75         3.78         0.36         5.60         11.75           14652         AAA2420         1.80         0.49         3.65         0.36         5.41         10.03           3576.28         AVA42420         1.81         0.60         3.65         5.62         0.55         0.68           146740         R77946         protatele	2	712950	AA282273	1-08	38.22	â	5.50	8.84	1.04	0.35	4.37	12.92	7
61331         T60111         4.13         0.82         5.03         11.18         1.30         0.80           287728         NS9158         3.49         0.71         4.90         14.39         0.35         0.46           301735         N80682         esophagus         4.10         0.80         4.56         9.05         2.75         2.83           811162         AAA86471         thyrold         28.87         6.85         4.21         0.70         3.01         6.33           36278         AAA001219         lymph         5.53         1.41         3.91         2.64         2.34         0.76           14008509         AA868629         borne         2.84         0.75         3.78         0.36         5.80         11.75           788953         AA42420         .         1.80         0.49         3.65         0.86         5.41         10.03           357628         W94121         .         1.80         0.49         3.65         0.55         0.68           146740         R77946         prostate         30.23         8.55         3.53         1.47         0.62         0.73           782335         AA432270         ardothselial <t< td=""><td>=</td><td>41889</td><td>R66438</td><th>adrenai</th><td><del>•</del></td><td>0.28</td><td>5.03</td><td>1.46</td><td>5.93</td><td>15.74</td><td>1.30</td><td>0.71</td><td>2</td></t<>	=	41889	R66438	adrenai	<del>•</del>	0.28	5.03	1.46	5.93	15.74	1.30	0.71	2
287728         N59158         3.49         0.71         4.90         14.39         0.35         0.46           301735         N90862         esophagus         4.10         0.90         4.56         9.05         2.75         2.93           811162         AAA86471         thyrold         28.87         6.85         4.21         0.70         3.61         6.33           38278         AAA86829         bone         2.84         0.75         3.78         0.36         5.80         11.75           788693         AAA2220         3.43         0.84         3.66         0.86         5.41         10.03           357626         WA4121         1.80         0.49         3.65         5.62         0.55         0.68           146740         R77946         prostate         30.23         8.56         3.53         1.47         0.62         0.73           782335         AA432270         endothselial         4.04         1.30         3.53         1.47         0.62         0.73	. 2	81331	T80111		4.13	0.82	5.03	11.18	1.30	0.80	8.59	3.28	7
301735         N90662         esophagus         4.10         0.80         4.56         9.05         2.75         2.83           811162         AAA86471         thyrold         28.87         6.85         4.21         0.70         3.61         6.33           38278         AAA001219         hymph         5.53         1.41         3.91         2.64         2.34         0.78           1406509         AAA86822         borne         2.84         0.75         3.78         0.36         5.80         11.75           768053         AAA2420         -         3.43         0.84         3.65         0.86         5.41         10.03           357628         W064121         -         1.81         0.60         3.65         5.62         0.55         0.68           195365         R88604         -         1.81         0.60         3.65         6.03         1.45         0.88           145740         R77946         prostate         30.23         8.55         3.53         1.47         0.62         0.73           782335         AA432270         endothrelial         4.04         1.30         3.12         6.21         5.84         1.80	<b>.</b> 5	287728	N59158		3.49	0.71	<b>4</b> .90	14.39	0.35	0.46	3.86	5.42	2
811162         AA486471         thyrold         28.87         6.65         4.21         0.70         3.61         6.33           36278         AA486471         thyrold         28.87         6.65         4.21         0.70         3.61         6.33           1406509         AA486829         borne         2.84         0.75         3.78         0.36         5.34         0.78           768633         AA424920         3.43         0.84         3.65         0.36         5.41         10.03           357628         WA4121         1.80         0.49         3.65         5.62         0.55         0.68           165740         R78964         1.81         0.60         3.62         6.03         1.45         0.88           145740         R77948         prostate         30.23         8.55         3.53         1.47         0.62         0.73           782335         AA432270         endothelial         4.04         1.30         3.12         6.21         5.84         1.80	<b>±</b>	301735	N90882	esophadus	4.10	0.80	4.56	9.05	2.75	2.83	0.89	7.09	7
382278         AA001219         lymph         5.53         1.41         3.91         2.64         2.34         0.78           1408509         AA888029         borne         2.84         0.75         3.78         0.36         5.80         11.75           788953         AA424920         3.43         0.84         3.65         0.86         5.41         10.03           357628         WA04121         1.81         0.60         3.65         5.62         0.55         0.68           185040         1.81         0.60         3.62         6.03         1.45         0.88           145740         R77948         prostate         30.23         8.55         3.53         1.47         0.62         0.73           782335         AA432270         endothelial         4.04         1.30         3.12         6.21         5.84         1.80	5	811162	AA486471	thyrold	28.87	6.85	4.21	0.70	3.61	6.33	4.17	8.28	7
1408509         AA424920         2.84         0.75         3.78         0.36         5.80         11.75           788953         AA424920         3.43         0.84         3.66         0.86         5.41         10.03           357628         W64121         1.80         0.49         3.65         5.62         0.55         0.68           195365         R88904         1.81         0.60         3.62         6.03         1.45         0.88           145740         R77948         prostate         30.23         8.55         3.53         1.47         0.62         0.73           782335         AA432270         endothelial         4.04         1.30         3.12         6.21         5.84         1.80	5	382278	AA001219	Manny	5.53	1.41	3.91	2.64	2.34	87.0	6.56 50	7.23	8
788953         AA424920         3.43         0.84         3.65         0.86         5.41         10.03           357628         W64121         1.80         0.49         3.65         5.62         0.55         0.68           195365         R88904         1.81         0.60         3.62         6.03         1.45         0.88           145740         R77948         prostate         30.23         8.55         3.53         1.47         0.62         0.73           782335         AA432270         endothelial         4.04         1.30         3.12         6.21         5.84         1.80	4	1409509	AA868929	poue	75.	0.75	3.78	0.36	5.80	11.75	0.59	0.31	2
357626         W04121         1.80         0.49         3.65         5.62         0.55         0.68           196365         R88604         1.81         0.60         3.62         6.03         1.45         0.88           14570         R77946         prostate         30.23         8.55         3.53         1.47         0.62         0.73           782335         AA432270         endothseltal         4.04         1.30         3.12         6.21         5.84         1.80	<b>5</b>	786953	AA424920		3.43	200	3.66	0.96	5.41	10.03	0.83	0.97	7
196365         R88904         1.81         0.60         3.62         6.03         1.45         0.88           146740         R77948         prostate         30.23         8.55         3.53         1.47         0.62         0.73           782335         AA432270         endothelial         4.04         1.30         3.12         6.21         5.84         1.80	9	357628	W94121		1.80	0.40	3.65	5.62	0.55	0.68	2.00	9.41	2
145740 R77946 prostate 30.23 8.55 3.53 1.47 0.62 0.73 782335 AA432270 endothelial 4.04 1.30 3.12 6.21 5.84 1.80	8	195365	R88904		1.81	0.60	3.62	6.03	1.45	0.88	3.51	6.23	2
782335 AA432270 endothelial 4.04 1.30 3.12 6.21 5.84 1.80	2	145740	R77948	prostate	30.23	8.55	3.53	1,47	0.62	57.0	5.51	9.44	7
	a	782335	AA432270	endothellal	<del>*</del> .04	1.30	3.12	6.21	5.84	1.80	0.66	1.08	77

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostastic Epithelial Cell Line: Genes upregulated fold or higher in 3 of 3, 5 fold or higher in at least 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.

Order	IMAGE Clone ID	Gen Bank Accession Number	MPM1(PrEC)	Ave Cancer Ceti Lines Values	Fold(AveCell line/MPM1)	Fold(MP <i>M2/M</i> PM1)	Fold(AveCell Fold(NPMZ/M Fold(MPMZ71/M F. Ins/MPM1) PM1)	PM1)	PM1) Cell Lines
			į	į	£01.67	7.33	66.37	1701.31	ю
-	757165	AA443950	0.01		0.100	7 27	944.19	9.33	ო
~	510631	AA099431	0.01	3.19	518.50	5	680.68	14.13	က
m	592707	AA160606	0.01	1.66	234.30	9 6	570.75	10.82	m
•	220851	H95633	0.01	1.28	196.56	8. 10.	2000	7 16	er.
	TROBOA	A A 4 4 B 1 B 6	0.01	1.81	180.70	8.88	526.08	2 6	) r
n (	10000	0.000000		2.58	117.17	3.60	344.82	3.10	<b>,</b>
ю	20029			1 12	112.48	13.45	301.13	<b>2</b> 2.83	n
_	1343732	•	5 6	2	103.67	3.09	302.09	5.84	m
∞	768226	AA424900	50	5	9	27.4	290.43	3.41	က
ø	731048	AA421275	0.01	0.99	<u> </u>	, t	249.63	3.45	೮
9	366039	AA071503	0.03	2.66	20.02	900	730.88	12.54	6
=	795627	AA459917	0.01	0.51	85.17		233.00	7 74	er.
: ;	4484830	•	0.01	0.71	71.26	3.19	203.18	• •	<b>,</b>
2	201011		£0.0	2.20	71.17	3.28	205.85	4.38	9 1
t	17515/	A4421047	3 6	990	65.30	45.84	106.60	43.45	מי
7	784296	AA447079	50	9 6	13 63	16 10	138.91	35.70	က
5	196569	AA460463	0.01	0.35	93.37	r.	168.10	11.76	m
5	346942	W94289	0.01	0.61	02.0	5.6	170.43	4.30	n
17	795858	AA461522	0.03	1.60	60.34 4.03	RY S	2501	70.45	m
<b>. .</b>	131452		0.45	26.93	59.79	104.24	90. i	47.07	, r.
2 \$	1558655		0.08	4.36	54.32	97.06	17.92	in: 1	) <b>r</b>
9 6	743018		0,01	0.50	50.47	15.70	123.80	3 :	, (
3 7	2077	•	0.10	5.02	49.93	99.52	44.77	9.4°	, c
7	71400		O SO	36.21	45.35	56.36	29.19	50.51	,
23	330/6			41.6	45.16	91.62	20.97	22.90	ო
R	626390	•		2.5		25.03	101.07	6.93	ø
24	128083	R09747	0.03	<b>?</b> *	\$ ! \$ !	2000	120.65	6.23	က
25	1389018	AA855158	0.05	0.81	43.48	7 (F	424.33	3.16	m
79	28498	R37395	0.01	0.43	43.37	20.0	400 63	50.5	177
27	757368	•	0.01	0.42	41.72	4.59	109.33	4 97	. 67
28	813633	_	9.0	1.68	41.51	3.55	10.00	ì	)
ì					•				

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least

0.01         0.41         41.11         4.81         99.20           0.01         0.38         38.18         3.08         106.34           0.01         0.30         37.18         16.25         80.78           0.01         0.32         38.78         12.25         87.45           0.01         0.32         35.62         5.91         86.27           0.01         0.25         34.44         20.19         73.72           0.02         0.33         33.18         5.81         82.72           0.01         0.33         33.44         20.19         73.72           0.02         0.59         32.57         29.44         39.40           0.03         33.18         5.81         82.77         30.24           0.04         0.32         31.52         8.48         9.21           0.05         0.24         29.16         9.10         30.24           0.07         0.44         29.13         4.48         9.21           0.03         0.44         29.48         5.18         9.20           0.04         29.24         29.48         5.18         5.03           0.04         20.23         25.24

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

W90036         0.01         0.21         20.90         5           HZ2824         0.01         0.25         20.42         16           HZ2653         0.01         0.20         20.40         5           HZ563         0.01         0.20         20.40         5           HZ54036         0.01         0.17         18.85         3           AA083187         0.02         0.31         18.24         16           AA087195         0.02         0.31         18.24         16           AA04741         0.17         2.95         17.83         7           AA047441         0.17         2.95         17.83         7           AA0476596         0.01         0.21         17.25         22           AA0476996         0.01         0.02         0.04         0.63         17.15           N867699         0.01         0.04         0.63         17.15         16.34           N867699
W90036         0.01         0.21         20.90         5           H22824         0.01         0.25         20.42         18           H22824         0.01         0.25         20.42         18           H22653         0.01         0.20         20.40         5           H2563         0.01         0.02         20.40         5           R5492         0.01         0.17         18.85         3           AA04741         0.01         0.17         18.85         3           AA04741         0.17         2.95         18.01         4           AA04741         0.17         2.95         17.19         7           AA04741         0.17         2.95         17.19         7           AA04355         0.01         0.01         0.21         17.35         2           AA609304         0.05         0.86         17.19         7         16.89           AA576996         0.01         0.01         0.17         16.89         17.19           NW81432         0.01         0.04         0.63         17.15         17.36           NW81432         0.01         0.04         0.63         17.15
W90036         0.01         0.21           W90036         0.01         0.25           N32542         0.01         0.25           N32542         0.01         0.20           H22663         0.12         2.46           R54036         0.01         0.19           R54492         0.01         0.17           AA883167         0.02         0.31           AA067441         0.17         2.95           AA701315         0.01         0.21           T49355         1.96         33.78           AA608304         0.05         0.86           NA7012         0.01         0.21           NA81432         0.01         0.17           NB1554         0.04         0.63           AA41846         0.01         0.16           H79466         0.07         1.17           H81554         0.07         1.17           H81554         0.07         1.17           H81546         0.02         0.32           AA41846         0.02         0.32           AA41846         0.02         0.32           AA406061         0.05         0.70           AA40606
W90036         0.01           HZ2824         0.01           N32542         0.01           HZ2663         0.12           R54036         0.01           R54036         0.01           AA883187         0.02           AA6047441         0.07           AA70315         0.01           AA606304         0.05           NA7012         0.01           AA606304         0.05           NA606304         0.01           AA606309         0.01           NW81432         0.01           NW814546         0.07           AA418546         0.02           AA404175         0.02           AA404175         0.02           AA406661         0.05           AA406661         0.05           AA406661         0.02           AA406641         0.01
W90036 W90036 H22824 N32542 H22863 R54036 R54492 A883187 A067195 AA057195 AA04741 AA701315 T49355 AA608304 N47012 AA192784 AA608304 N81432 N26769 W81432 N26769 W81432 AA41846 AA045175 AA418466 AA045175 AA41846 AA406061 R40434
451581 40010 39722 467161 488431 435099 67440 031568 627521 454083 347686 259606 259606 259606 259606 47416 47416 47416 47416 47416 47416 47416 47416 47416

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upreguizted 3 fold or higher in 3 of 3, or 5 fold or higher in at least

99	1031446	AA609189	9.0	0.50	14.14	5.44	31.74	5.25	ဗ
_	32109	R42780	0.0	1.25	14.14	31.21	5.86	5.35	ო
	856289	AA774665	1.24	17.58	14.13	5.00	9.24	28.16	က
	30850	R42600	0.03	0.39	13.78	6.39	29.82	5.12	က
	41408	R56149	0.0	0.13	13.76	17.85	18.97	4.48	ო
\$	035796	AA628867	4.1	15.63	13.71	10.64	9.76	20.74	က
	898286	AA598974	1.21	16.52	13.65	13.47	11.86	15.62	က
	502003	AA128617	0.01	0.17	13.58	7.33	15.99	17.41	6
	359269	AA016225	0.0	0.13	13.43	9.24	8.92	22.12	က
	415089	W93379	0.65	8.63	13.24	8.23	12.04	19.45	ო
	1456409	AA885478	0.14	1.91	13.23	22.16	9.42	8.12	٣
	824074	AA491227	0.02	0.26	13.22	3.39	18.08	18.18	က
	51103	H19217	0.03	0.23	13.14	14.72	17.85	6.86	ო
	1467166	AA883181	0.08	1.05	13.14	3.08	29.16	7.18	က
	244050	N34042	0.01	0.17	12.88	5.65	21.72	11.28	က
114 10	1055121	AA621355	20:0	0.25	12.67	6.31	28.51	3.21	es
	897807	AA598531	0.29	3.56	12.37	21.96	7.81	7.37	က
116 8	845441	AA644550	0.0	0.12	12.30	5.26	7.58	24.06	ო
	686172	AA262211	1.18	14.39	12.18	5.65	13.80	17.09	ო
118 10	048079	AA778717	9.0	0.31	12.10	3.28	15.47	17.55	m
119 78	785368	AA476578	85.	16.58	12.06	8.77	4.23	23.17	က
120	451649	AA706901	04.	16.82	11.98	4.34	18.70	12.88	n
	46461	H08940	0.33	3.89	11.80	3.09	13.18	19.11	n
•	030798	AA609002	0.03	0.31	11.73	10.15	8.33	16.72	ო
	33839	R44816	96.0	11.24	11.67	3.60	19.31	12.09	က
	357138	W83523	0.02	0.20	11.63	10.87	20.61	3.40	ო
125 36	366971	AA026682	2.38	27.64	11.61	3.65	10.18	20.99	ო
_	809228	AA179161	0.01	0.11	11.46	4.58	5.70	24.11	က
127 21	210820	H67712	9.04	0.45	11.30	5.06	7.68	20.94	6
	795746	AA460299	0.52	5.68	10.87	17.71	9.89	<b>2</b> 6.	ы
129 79	95840	AA461513	0.01	90.0	10.81	4.64	15.78	12.02	က
_	88209	AA453433	0.01	60.0	10.70	11.07	8.10	12.92	ო
131 76	67403	AA417920	0.56	5.93	10.66	4.43	22.31	5.22	6
132 20	04483	H58234	1.27	13.51	10.64	11.48	13.18	7.28	က
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Table 2-10; Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line. Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.

ო	m	က	က	m	က	n	ო	က	ო	n	က	က	က	6	က	က	m	က	က	က	ო	က	က	က	က	က	ო	က	ო	e	3	e	က	က	
5.03	4.82	22.97	5.78	14.06	4.93	16.03	15.10	6.87	14.13	18.44	7.37	3.91	5.79	11.50	7.50	9.62	18.79	15,70	8.67	7.35	5.24	4.33	11.30	28.2 28.	4.46	13.62	5.71	13.08	6.90	10.11	11.70	5.53	15.48	<b>78</b> .6	
5.77	22.27	4.07	20.48	3.07	3.47	9.94	8.56	18.90	9.81	6.58	10.28	20.10	10.79	11.42	14.40	13.22	5.97	7.97	15.51	11.24	12.44	10.85	11.63	8.66	14.83	8.45	8.47	10.73	12.47	7.21	9.17	13.78	6.65	10.99	
20.21	3.65	3.39	3.74	12.71	21.35	3.40	5.69	3.45	4.99	3.83	11.04	4.57	11.98	5.53	6.47	5.48	3.48	4.56	3.79	9.36	10.06	12.49	4.58	12.99	8.17	5.38	13.21	3.46	7.38	9.14	5.30	69.9	3.86	5.02	
10.34	10.24	10.15	10.00	9.95	9.92	9.79	9.78	9.74	9.64	9.62	9.57	9.53	9.52	9.48	9.46	9.44	9.41	9.41	9.32	9.32	9.24	9.22	9.17	9.16	9.15	9.15	9.13	60.6	8.92	8.82	8.72	8.67	8.66	8.62	2
2.7	0.10	0.18	0.37	1.05	5.07	0.57	0.32	12.26	0.17	0.10	0.16	31.33	10.61	1.40	2.04	10.53	7.14	5.85	0.19	3.43	1.03	0.97	9.89	0.60	3.21	4.	1.40	0.26	0.11	0.05	4.90	12.90	48.92	3.90	
0.74	0.01	0.02	700	0.11	0.51	0.06	003	1.26	0.02	0.0	0.02	3.29	1.12	0.15	0.22	1.12	0.76	0.62	0.02	0.37	0.11	0.10	1.08	0.07	0.35	0.16	0.15	0.03	0.01	0.01	0.56	1.49	5.65	0.45	
AA678295	AA626028	AA291558	A A 508787	AA625979	B44734	A1003636	AA041197	AA682533	AA157261	AA156342	AA701652	N33274	N47113	62208N	AA702684	N24042	AA678348	AA204830	W74377	AA045665	H04789	AA705047	AA446462	H77727	H10192	H46176	AA598817	R69584	W49672	T50699	AA187351	AA137072	H95638	AA520999	
430928	745019	725308	808073	745490	24124	391987	376452	431242	590500	590727	433567	273546	280375	303099	448059	269300	430973	645565	346552	488213	43733	461488	781047	234617	46694	177884	897966	141726	324901	78217	624627	491184	242952	826273	
2	<u> </u>	3 \$	3 5	<u> </u>	5 5	3 5	3 4	645	<u> </u>	<u> </u>	145	34.	147	148	149	150	15.	152	153	<u> </u>	155	156	157	158	159	160	161	162	163	20	165	166	167	168	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line. Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.

08.0	6.37	14.40	6.83	4.37	8.51	7.71	14.93	15.23	15.35	13.11	4.58	5.12	6.49	10.12	4.46	10.00	11.39	10.84	13.28	11.71	9.05	7.97	11.50	7.15	8.23	5.28	4.93	12.46	12.18	7.87	9.46	14.01	4.48
14.21	13.73	6.65	8.33	16.37	11.10	13.64	5.10	3.98	4.12	7.53	15.49	6.62	11.97	9.24	7.24	8.93	4.17	7.66	3.58	8.00	5.22	7.09	5.10	4.96	10.88	4.4	<b>3</b> 6.0	4.68	6.84	6.65	4.6	4.60	13.99
7.42	4.95	3.89	9.78	4.16	5.27	3.04	4.35	5.06	4.69	3.42	3.96	11.86	4.88	3.94	11.54	4.23	7.56	4.60	6.23	3.35	8.56	7.77	6.04	10.53	3.47	12.64	10.35	5.03	3.00	7.30	6.01	3.23	3.25
8.51	8.35	8.32	8.31	8.30	8.30	8.13	8.13	8.09	8.05	8.02	8.01	7.87	77.7	77.7	7.75	7.72	7.70	7.70	7.70	7.68	7.61	7.61	7.55	7.55	7.53	7.44	7.41	7.39	7.34	7.31	7.30	7.28	7.24
0.14	0.19	4.39	1.57	80.0	7.83	0.25	21.34	10.44	81.74	0.28	5.63	0.54	2.89	0.60	5.81	0.35	24.59	13.42	1.68	261.61	5.69	5.53	0.15	9.0	5.64	0.20	21.88	23.90	57.27	3.90	0.12	3.80	0.41
9.0	0.02	0.53	0.19	0.0	<b>3</b> 6.0	8	263	25	797	8	0.70	0.07	0.37	90.0	0.75	90.0	3.19	1.74	020	7.8	0.75	0.73	0.02	0.01	0.75	0.03	2.95	3.23	7.80	0.53	0.02	0.52	90:0
T52311	AA485265	AA455786	R60019	W73790	AA425755	W96216	P41329	HK47K9	T65902	AA705825	H29521	AA455938	H29897	AA625764	AA431741	AA906257	T96718	155607	W15305	AA676460	R96941	AA608568	N30372	AA070331	AA434403	T61122	AA701455	N83924	AA488367	H29227	AA477400	N70714	H16833
1877	340150	809557	42906	344134	773383	358599	3014B	2012	80399	292182	52741	813271	190325	745393	782259	505919	121251	73596	322461	882510	200402	069098	260035	530820	770879	83666	435076	309288	843163	52704	740620	294273	50787

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

,	m	n	es .	es.	E)	n	E.	n	က	<sub>6,0</sub>	n	n	n	· (*)	, ,	<b>,</b>	n	n	E.	က	m	က	m	e	6	6	က	e	ຕ	60	e	8	3	n	63	es	
	5.38	8.99	8.78	10.69	3.65	7.98	6.50	9.20	11.35	3.14	9.86	3.89	5.69	4 49	3 5	12.11	12.23	5.38	6.30	4.64	10.97	7.65	4.14	5.60	6.95	8.44	7.62	8.12	4.80	12.51	3.05	4.63	5.67	11.00	10.01	7.97	
	11.90	9.60	7,63	6.35	5.87	7.83	9.95	7.07	6.52	11.26	4.41	3.38	10.31	96.3	000	4.92	5.03	10.24	4.78	9.34	3.78	4.68	8.01	5.02	99.6	4.39	4.47	3.71	7.44	3.32	5.69	69.6	8.97	4.30	5.16	8.32	
	4.39	3.01	6.15	4.40	11.72	5.35	4.65	4.74	3.01	6.40	6.49	13.48	478	000	0.00	3.47	3.20	4.75	9.29	6.37	5.57	7.84	8.02	9.45	7.5	7.13	7.85	8.09	7.52	3.80	10.89	5.27	4.93	4.19	4.26	3.07	; ;
	7.22	7.20	7.18	7.15	7.08	7.05	7.03	7.00	90 9	693	26.9	50.5	20.0	26.9	6.85	6.83	6.82	6.79	6.79	6.79	6.77	6.72	6.72	69.4	90:0	9 <b>4</b>	5.65	6.64	6.59	45.5	45.9	6.53	6.52	05.50	0.0	6.45	} r
	33.83	6.75	18.03	0.10	14.78	3.05	0.08	100.32	470	2 0	9 09 2	54.45	61.13	7.46	0.72	2.56	8.14	0.08	0.34	8.93	27.70	030	3.5	50.0	10.7	20.00	10.13	7 2	100	35.62	0.42	7.31	2 64	פני נינ פני נינ	27.77	‡ 4c	07:1
	4.68	9	2.51	0.01	2.09	0.43	6	14.33	2	3 8	3 6	3 6	3. 6	<b>6</b>	0. 0.	0.37	1.19	0.01	0.05	1.32	4.09	2	2 2	ţ ;	; i	2 2	70.70	2 6	3 6	27 4	800	1.12	0.41		? f f		<u>2</u> .
	H97488	A A 4 B 0 9 5 2	AA663310	AA456611	N45236	Present	674203	A1046350	AU15558	W80180	K36046	NZMSOS	W72437	AA465238	AA701645	N57722	R23055	AA256231	H71230	HORSON	A A O 10065	H47430	A 6 500 733	AA609723	K2828/	AA465593	AA589092	M4430033	H33203	A 4 3 0 7 8 4 3	MOS645	H20847	N48075	2000	600/LH	HZ1343	AA045506
	251936	798114	853368	809567	283173	126650	143365	1637743	2000	00014	40773	704207	345525	814225	433576	246659	131316	681906	214604	46091	350110	2003	20010	0001001	20.00	814246	950445	2010	*12907	126.054	43667	51460	281733	3 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6	2020	2000	401531
	204	20,5	206	207	2 6	9 6	2 6	2 7	17	212	213	214	215	216	217	218	219	220	22	; ;	3 5	3 2	<b>5</b> 7 6	87	8	227	528	R77	75.0	3 8	7	22.6	3,5		87 5	757	857

culated 3 fold or higher in 3 of 3, or 5 fold or higher in at least Table 2-10: Prosta

state Ca	ncer Cell	Lines vs. No	mal Prosta 2 of 3 or 1	atic Epithelial 0 fold or high	Cell Line: Ge ıer in at least	mal Prostatic Epithelial Cell Line: Genes upregulated 3 fold 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines	ited 3 fold or h cell lines.	igher in 3 of 3, o	state Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or nigner in tate Cancer Cell lines.
200	430007	AA034115	0.26	1.70	6.43	4.03	5.78	9.48	ĸ
2 6	395955	AA757564	0.13	0.82	6.41	5.00	4.97	9.25	က <sup>(</sup>
74	782476	AA431753	0.76	4.82	6.38	8.52	6.40	<b>2</b>	rs (
242	343298	W68162	0.10	0.62	6.37	10.68	3.55	4.86	es :
243	151055	H02231	7.49	47.65	6.36	6.67	5.78	6.63	6
7	280907	N50827	0.07	0.47	6.32	3.73	12.06	3.15	en.
245	32050	R41972	2.0	2.79	6.31	4.19	3.05	11.67	က
246	782594	AA447531	<b>60</b>	9.0	6.30	3.58	10.41	4.91	<b>.</b>
247	292936	N63744	0.72	4.53	6.29	3.26	6.93	8.69	m
248	130004	R11605	0.38	2.35	6.25	3.19	6.05	10.52	6
2,49	22895	R38640	0.02	0.15	6.24	3.63	10.49	4.60	က
250	249603	H84871	5.09	31.76	6.24	11.55	3.15	4.01	m
251	796694	AA460685	6.55	40.75	6.23	4.27	10.33	4.07	r.
252	813586	AA447662	1.97	12.17	6.19	4.19	5.59	8.79	ന
25.	137890	R68581	0.76	4.70	6.17	4.68	3.86	96.6	ന
<u> </u>	128947	R10284	1.20	7.33	6.13	6.92	5.67	5.79	60
255	50983	H18017	0.02	0.14	6.12	7.62	5.90	4.83	m
256	742115	AA405800	0.67	3.47	6.11	8.6	4.21	4.48	ю
257	773073	AA425302	2.73	16.66	6.10	7.47	5.73	5.08	ю
258	256680	H96392	0.81	4.93	90'9	3.55	6.77	7.86	m
259	489805	AA102068	0.01	0.08	90.9	6.43	4.55	7.19	က
260	814306	AA459318	.89	29.59	90.9	6.00	6.93	5.21	n
261	773147	AA425404	1.31	7.89	6.04	4.66	8.09	5.37	က
262	951068	AA620437	0.16	0.94	6.02	3.62	3.03	5.40	6
263	275871	R93875	0.01	0.07	9.69	5.03	7.41	5.52	က
797	271078	N29918	25.0	3.18	5.90	7.03	3.15	7.53	က
265	301104	N81049	0.24	1.38	5.88	9.69	3.97	3.97	က
268	813827	AA453714	0.02	0.10	5.87	3.43	4.55	9.64	<b>6</b>
267	46154	H09065	0.01	90.0	5.87	3.67	4.93	9.01	r
268	489594	AA099523	0.37	2.15	5.86	4.97	9.49	3.11	n
209	357531	W94009	1.37	8.00	5.86	4.55	5.43	7.60	ю
270	44255	H06113	222	12.98	5.83	3.48	4.46	9.58	ю
27.1	809535	AA454585	7.47	43.39	5.81	3.50	5.22	871	က
212	41188	R56870	2.45	14.21	5.81	3.49	4.78	9.15	е,
273	287637	N59136	1.87	10.85	5.80	4.53	5.30	7.58	က
					<b>&amp;</b>				

Table 2-10: Prostate

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithellal Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

m	၈	ო	ന	6	က	ო	m	၈	က	en .	6	<b>6</b> 0	က	၈	က	ဂ		n	က	<b>6</b> 0	ო	೮	•	က	e	6	က	ဗ	က	٣	9	က	m	ы	
6.97	5.45	5.33	4.75	8.15	4.97	4.67	5.63	4.55	¥.4	625	4.80	5.33	5.26	4.03	6.82	5.61	6.41	5.90	4.80	5.64	6.54	4.37	6.23	4.69	3.60	5.29	3.08	3.05	3.31	6.55	5.01	20.0	3.51	4.93	
5.27	9.00	5.12	4.23	3.76	3.23	7.05	5.03	6.00	4.38	4.66	5.78	99.9	3.76	5.47	4.74	5.83	3.84	5.34	4.57	5.49	4.24	4.27	4.29	3.94	3.35	4.33	7.24	4.15	4.89	3.95	3.86	5.23	7.07	4.94	
3.25	4.99	4.34	6.39	3.45	7.16	3.62	4.62	4.64	6.45	4.26	4.49	3.05	6.00	5.36	3.28	3.32	4.46	3.38	5.12	3.32	3.64	5.78	3.85	29.62	7.29	4.52	3.82	6.91	5.91	3.56	5.16	5.09	3.32	<b>4</b> .03	
5.16	5.15	5.13	5.12	5.12	5.12	5.11	5.09	5.06	5.06	9.09	5.02	5.02	5.01	4.96	4.95	4.92	4.90	4.87	4.83	4.81	4.81	4.81	4.79	4.75	4.75	4.71	4.71	4.70	4.70	4.69	4.67	4.65	4.63	4.63	10
2.47	5.19	31.82	0.40	0.24	34.01	2.84	0.77	10.81	23.18	8.96	1.02	23.47	4.55	0.75	24.26	11.53	6.70	5.68	<u> </u>	0.90	0.81	15.65	1.22	4.83	5.71	0.45	3.94	32.63	3.65	0.19	3.48	1.69	3.36	0.43	
0.48	1.01	6.20	80.0	0.05	9.0	0.58	0.15	2.14	4.58	1.7	0.20	4.68	0.91	0.15	4.90	23.	1.37	25.0	0.32	0.19	0.17	3.26	0.25	1.02	1.20	0.09	98.0	6.94	0.78	8	0.74	9.36	57.0	0.09	
AA136060	AA703000	N66132	R80387	AA663995	AA450265	H18068	AA885851	AA463924	N93715	AA448160	AA683557	AA504272	AA129552	AA626784	N40939	AA443969	AA459532	AA418918	AA678135	AA156599	AA776692	AA620553	AA481757	N90523	N50745	AA883127	AA621019	AA448396	N73301	H97000	T66936	AA046321	H23482	AA682405	
490060	447208	278483	147147	855390	789182	50849	1500388	810264	307119	782766	505864	825411	564803	877767	277163	757206	810974	767994	431895	502296	1292828	951142	810785	306146	283751	1467026	1056200	781341	292068	249949	90499	376764	52543	461775	
308	340	2 5	3.5	3 5	314	315	316	317		319	320	i s	322	323	70.	7.	32, 52	327	328	329	330	33.	332	333	33.	335	336	337	338	339	340	**	342	85	

pregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least Table 2-10: Pro

;	KAAEEA	AANZAEER	4.89	22.62	4.63	3.06	3.22	7.60	ო
<b>,</b>	505400	A 4 173926	2.40	11.07	4.60	3.28	5.08	5,45	ന
2 2	202033	N60491	0.67	3.08	4.60	4.85	3.14	5.80	က
	287749	N62245	177	8.14	4.60	4.35	3.56	5.89	က
<b>3</b> 3	SE1336	A A 600762	0.22	66.0	4.58	6.38	4.05	3.34	က
\$ 3	306684	NITABEE	15.03	68.72	4.57	3.02	5.75	4.95	m
2	100042	N/4800	0 0	9.0	4.56	5.14	4.81	3.75	က
3	1000	145517	£ 2 4	25.23	4.56	4.05	3.27	6.37	6
	00000	**************************************	4 25	292	4.55	4.56	3.51	5.59	က
3 5	001087	WCC830	2 70	16.81	45.54	3.79	4.39	5.44	ო
3	07/707	N30078		13.59	4.53	3.16	5.08	5.34	က
t i	500470	AA058305	4.33	19,58	4.52	3.27	6.13	<b>4</b> .18	က
3 3	2000	4470704	5	0.60	4.52	3.14	3.41	7.01	က
g i	10014	A 1017	 	3.85	15.51	5.43	3.36	4.75	ო
i s	02/50	AACOODE	29.84	133.80	4.48	3.35	5.80	4.30	ຕ
9 6	186016	A 4 6 4 0 4 7	2.35	10.51	4.48	5.20	5.13	3.12	က
B 6	400548	AA128700	1.22	5.43	4.46	5.70	4.50	3.19	m
3 5	795371	AA453273	2.35	10.45	4.45	4.16	5.42	3.76	ო
3 2	358457	W96114	7.33	32.61	4.45	3.43	5.36	4.54	m
3 5	35,6732	W84486	0.67	2.96	4.43	4.85	4.4	9.4	က
3 3	1627623	A1014704	0.49	2.19	4.43	5.61	3.81	3.87	ო
a a	2,1942	VA60015	2.28	10.0t	4.40	3.32	4.59	5.27	m
3 8	842906	AAABGA30	2	4.57	4.39	3.03	3.33	6.82	က
2 5	1475463	AA857804	1.80	7.88	4.39	5.77	3.14	4.26	က
9 6	350135	AA010128	0.47	2.08	4.38	3.23	3.76	6.16	က
	108174	P04833	23	0.93	4.38	4.72	3.33	5.10	ю
2 5	10000	200767	2	0.18	4.37	3.06	5.05	4.99	က
2	200642	NOTATE	2 20	9	4.36	5.17	4.31	3.59	က
5 6	324313	AA284108	0 0	0.09	4.34	3.11	6.72	3.19	က
	280342	N49261	4.13	17.91	4.33	3.32	4.21	5.47	m
374	40881	R56055	0.15	0.64	4.32	4.06	4.72	4.18	က
375	250519	H89698	1.26	5.41	4.31	4.10	3.51	6.33	ന
376	418150	W90164	0.62	2.65	4.31	3.30	6.39	3.24	က
377	258761	N30185	1.26	5.44	4.30	4.24	3.24	5.43	က
378	25274	R12804	0.39	1.66	4.29	3.40	5.79	3.69	က
					=				

al Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least Table 2-10: Prost

state Ca	incer Cell	Lines vs. Nor	nal Prostati 2 of 3 or 10	ic Epithelial ( fold or high	Cell Line: Ger er in at least	mal Prostatic Epithelial Cell Line: Genes upregulated 3 told 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines	ted 3 told of ni cell lines.	state Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold of nigher in 3 of 3, or 2, or 2, or 2, or 2, or 2, or 2, or 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.	
ŗ	70763	8081808	0.63	2.72	4.29	3.87	4.70	4.31	9
A 6	417855	W88702	1.03	8.22	4.27	3.03	4.54	5.24	e
200	1666058	A A D 7 5 3 8 8	0.19	0.82	4.25	3.90	3.93	4.93	၈
28.5	502767	AA126259	0.42	1.80	4.25	3.90	3.05	6.79	ю
705	453599	AA678569	1.66	7.01	4.23	9.00	3.07	3.61	ю
8 8	42793	R59722	0.30	1.26	4.22	3.06	5.07	4.55	ဗ
, K	183771	H44448	238	8.98	4.22	3.65	5.59	3.43	က
S 8	365157	AA024637	233	9.83	7.7	3.26	5.64	3.75	m
200	324255	W47254	1.68	7.07	4.21	3.45	4.51	4.67	က
\$ \$	809421	AA459909	3.57	15.01	4.20	3.52	6.97	3.12	ю
2 2	101867	N92478	0.36	1.52	4.20	3.30	4.63	4.67	m
	825089	48489247	0.11	0.45	4.19	3.43	4.75	4.40	က
9 6	165641	AA025937	1.29	5.38	4.19	3.00	3.59	5.97	6
	78198	Troate	2.26	9.46	4.18	3.30	5.98	3.26	e
766	788566	AA452966	1.36	99.9	4.18	3.56	3.76	5.22	က
8 8	755301	AA496360	0.05	0.22	4.16	3.57	3.72	5.18	က
295	1387760	AA838691	1.66	6.85	4.14	4.66	4.62	3.14	m
30, 20	510170	AA053129	6.42	26.51	4.13	3.38	4.00	5.01	ю
397	127925	R08935	0.28	1.14	4.13	3.18	4.98	4.22	n
388	845380	AA644099	1.43	5.90	4.12	4.14	4.91	3.31	<b>ю</b>
300	305538	N89861	5.14	21.16	4.11	4.96	4.11	3.26	က
907	648693	AA125825	0.49	2.02	4.11	4.24	3.39	4.71	en.
107	125183	R05693	7.53	30.82	4.09	3.66	3.99	4.63	m
<b>4</b> 02	289945	N64617	0.64	2.60	<b>4</b> .08	3.25	3.58	5.40	m
1 6	431803	AA678021	17.87	72.78	4.07	3.72	5.27	3.23	က
2	824739	AA489000	1.22	4.98	4.07	3.33	4.59	4.30	က
405	136180	R33303	5.09	8.49	4.07	3.09	4.10	5.02	es e
907	51469	H23985	2.11	8.55	4.05	3.08	3.13	5.93	ന
<b>4</b> 0 4	461592	AA705142	2.49	10.05	2.4	5.90	3.14	3.07	9
408	290378	N64508	0.80	3.22	4.03	3.72	3.29	5.07	က
409	795830	AA460571	90.0	٠. ج	4.00	3.83	4.74	4 ·	м
410	416390	W86860	2.20	9.11	3.99	3.72	3.60	4.63	က
=	150135	H01926	4.15	16.49	3.97	3.31	4.75	3.86	9
412	47681	H11792	8.31	32.97	3.97	3.15	3.42	5.33	ы
413	271006	N29901	3.83	15.20	3.96	3.30	5.01	3.59	m
					12				

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

	625764	AA188366	0.91	3.60	3.96	4	ŗ		•
7 4	1554974	A1023724	25	8.02	3.94	3.65	3.46	4.73	,
2 9	407305	HSORKE	0.29	1.14	3.93	4.53	3.70	3.55	7
9 !	20070	1130000 113306	35	1.32	3.93	3,19	4.45	4.15	ന
- 9	10003	0 0 4 4 9 8 7 8	7.7.	8.89	3.93	4.14	4.07	3.57	<b></b>
9 5	411128	AA211448	0.43	1.69	3.92	4.47	3.83	3.47	(F)
a 9	201210	NED 283	4.34	16.90	3.92	3.28	3.75	4.72	e-2
27	010007	202507		60.0	3.86	4.33	3.12	4.12	(7)
129	115230	19697/	30.5	11.74	3.85	5.38	3.09	3.09	~,
7	04500	AA436132	2,68	10.26	3.85	4.86	3.02	3.67	<b>(*)</b>
3	# 7 / / ·	BC10C1	3 6	11.13	3.84	4.81	3.36	3.36	~,
<b>\$</b> 7.	0.505.0	AM33036		4.47	3,82	4.55	3.33	3.58	(,)
<b>425</b>	295/10	/697/N		7	3.74	3.23	4.39	3,59	• •
426	884690	AA630016	5 6	5 6	3.74	3.07	3.48	4.65	••
427	377884	<b>AA777</b> 092	64.0	25.0	3.68	3.83	4.03	3.19	
<b>428</b>	453642	AA775888	3 i	\$ 5.00 P	3 69	4.08	3.46	3,51	• •
2	1586340	AA974348	2.83 0.00 0.00 0.00 0.00 0.00 0.00 0.00 0	10.01	9 89	. E.	3.85	3.86	••
8	757404	AA426341	5 G	\$.07 62 6	9 9	3.24	3.30	4.45	•••
<del>1</del> 31	745286	AA625558	2 :		3.66	3.28	4.40	3.28	۷,
<b>432</b>	773599	AA429422	<b>1</b>	9.00	99	3.48	4.23	3.25	~,
53	487723	AA043552	76.1	5 6	3 65	474	3.21	3.02	~,
3	288658	N79353	0.26	6.3	 	4 05	3.12	3.76	(*)
<u>გ</u>	279920	N57526	200	157	39.	3.48	4.25	3.19	~,
28	813648	AA453679	بر جون جون	1 2 2	1 6	3.72	3.26	4.37	••
437	136744	R35230	8 1	77.7	3.02	3.70	3.44	3.70	••
8	487426	AA046523	77.7	10.00	- u	96 r	3.72	3.07	•••
65	589479	AA157129	77	10.	5 4	50 E	3.86	3.62	•
₹	256260	H94617	<del>?</del> :		t (1	) r	3.73	3.24	
<u>₹</u>	827273	AA191461	5.	# 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10:00 10 10 10 10 10 10 10 10 10 10 10 10 1	30.0		78	3,32	•••
7	1604596	AA996230	0.80	2.80	94.0 94.1	67.6	3.6	3.59	•
3	843224	AA488445	5.65	19.64	3.47	, i.	3 .	Ş	•
3	781339	AA448394	0.19	9.64	3.42	3.04 4	4.21	3.06	
45	275653	R93309	0.72	2.45	3,41	3.42	3.22		
2 3	840384	AA485773	5.81	19.81	3.41	3.01	3.43	3.80	.,
ָרָ בְּ	442049	A A 707840	4 07	3.62	3.39	3.46	3.12	3.59	
Ì 3	27070	AA421273	1.36	4.57	3.37	3.82	3.16	3.13	••
		,							

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in at least

	60
	3.05
cell lines.	3.55
2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.	3.08
er in at least 1	3.23
0 fold or high	30
2 of 3 or 1	4

501731	AA127861	2.15	6.95	3.23	3.08	3.65	3.05 3.17	ი ო
ATA9	5 5	0.0	9.92	991.80	1.38	2966.06	2.96	2
AA009615	515	0.0	8.39	838.64	1.96	2508.90	7.05	7
R609	98	0.01	9.14	660.46	2.32	61.14	1917.92	<b>7</b>
N57483	63	0.01	9.00	613.07	1.58	1831.96	5.67	7
A433	885	0.0	5.08	508.28	2.41	1516.35	<b>G</b> .10	7 (
T973	3	9,0	18.56	445.85	3.24	1331.89	242	7
<b>₩</b> 78	108	0.01	4.34	433.98	2.48	1294.73	4.72	7
N N	379	0.01	<b>3</b> .0	393.88	0.73	1171.33	9.57	7
AAA	706	0.01	2.83	283.26	1.00	841.07	7.73	2
AAAA	848 848	0.01	1.70	228.86	10.58	674.66	1.35	7
AAA	6866	100	1.58	157.55	19.1	467.71	3.31	7
AA	9292	0.01	<u>.</u>	154.10	3.95	457.34	1.00	61
₹ ₹	59154	0.02	2.74	114.55	1.41	338.20	4.03	7
Ĩ	533	0.02	1.93	95.78	0.80	140.40	146.14	7
¥	5214	0.08	6.83	85.65	0.89	150.46	105.59	7
X	611	0.01	0.82	81.58	1.42	229.57	13.76	7
A A	6833	0.01	0.71	70.86	2.42	205.80	4.35	7
₹	9486	0.08	5.16	65.49	0.87	95.72	99.88	7
*	1754	0.01	0.83	60.41	6.88	173.62	0.73	7
*	1784	0.05	2.87	58.88	1.03	51.92	123.70	7
*	4	0.22	12.41	26.02	78.84	88.50	0.72	7
Æ	AA086475	0.01	0.28	53.90	5.36	154.39	1.93	7
\$	123	9.0	2.21	52.11	0.54	150.40	6.38	7
¥8	73635	90.0	3.95	51.39	1.05	149.73	3.40	7
A	17224	1.20	60,94	50.82	144.61	1.37	5.47	7
₹ ¥	AA443938	0.19	9.88	50.76	4.68	145.07	2.54	7
3	741	0.02	1.10	50.72	0.63	139.36	12.17	7
AASC	1844	0.26	13.02	49.67	0.91	56.58	91.51	7
AAA	3802	0.01	0.49	49.10	3.38	142.92	1.00	2
AA90	988	0.26	12.64	47.88	137.81	27.2	3.10	7
\$	9120	0.01	0.60	45.99	2.24	128.30	7.45	7
85	R59936	0.0	0.46	45.56	1.92	130.70	4.06	7
\$	W292429	0.02	0.87	44.45	4.84	125.65	2.84	7
				4				

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

743562         AA400365         0.04         1.89           743562         AA400365         0.04         1.89           28422         R14230         0.46         17.33           280608         AA071089         0.01         0.38           774446         AA446120         0.05         1.69           428828         AA005254         0.01         0.35           428828         AA005254         0.01         0.35           196544         R91566         0.05         1.69           52928         AA45507         2.25         8.56           52928         H29290         0.08         2.62           74318         AA401434         0.02         0.70           815861         AA485052         0.26         7.89           489656         AA485052         0.26         7.89           489656         AA485052         0.02         0.70           53039         R15740         0.01         0.27           771294         AA460274         0.01         0.27           771294         AA460274         0.01         0.25           280699         N47445         0.65         0.39           2806	43.31			3	4
AA400365 AA867748 R14230 AA01629 AA46120 AA005254 0.01 AA159179 AA159179 AA425307 AA01634 AA485052 AA093369 AA401434 AA485052 AA485052 AA485052 AA485054 AA485054 AA485057 AA43624 AA45158 U.01 AA443624 AA45158 U.01 AA443624 AA45158 U.01 AA44364 U.01 AA701996 AA90948	37.79	2.77	123.35	3.82	64
AA867748  R14230  AA071089  AA446120  AA06254  AA159179  AA425307  AA425307  AA485052  AA485052  AA485052  AA485053  AA485054  AA485054  AA485057  AA485057  AA485057  AA485057  AA436274  O.01  AA436274  AA436274  O.01  AA436274  AA436274  O.01  AA443627  AA45158  O.07  AA701996  R15813  W94486  U.01  AA49045	•	5.72	105.49	2.18	7
R14230 AA071089 AA446120 AA466120 AA006254 B91566 AA159179 AA425307 AA425307 AA485052 AA485052 AA485052 AA485052 AA485053 AA485054 AA485054 AA485055 AA485055 AA425158 CO1 AA425158 CO1 AA425158 CO1 AA425158 CO1 AA701996 CO2 AA701996 CO2 AA701996 A19826 CO2 AA49045 CO1 AA490945 CO1 AA490945 CO1	CY 12	0.83	107.04	4.40	2
AA071089 0.01 AA446120 0.05 AA06254 0.01 R91566 0.25 AA425307 2.25 R59473 0.03 H29280 0.08 AA401434 0.02 AA485052 0.26 AA099369 0.01 R49439 0.21 R49439 0.21 R49439 0.01 AA460274 0.01 AA46055 0.63 AA425158 0.01 H50655 0.63 AA425158 0.01 AA701996 0.07 AA701996 0.07 AA701996 0.01	36.73	3 80	103.70	1.11	2
AA446120 AA46524 R91566 AA159179 AA425307 R59473 R59473 R49439 R49436274 R4018 R49439 R4018 R4018 R4018 R4018 R4018 R4018 R4018 R15813 R9647 R15813 AA49045 R496647 R15813 AA490945 R331	36.23	1 20	103.43	40.4	2
AA005254 0.01 R91566 0.25 AA159179 0.18 AA425307 2.25 R59473 0.03 H29290 0.03 AA401434 0.02 AA401434 0.02 AA405052 0.26 AA405052 0.26 AA405054 0.01 AA460274 0.01 AA443624 0.01 AA443624 0.01 AA425158 0.01 AA425158 0.01 AA425158 0.01 AA425158 0.01 AA425158 0.01 AA401996 0.07 AA701996 0.01	35.22	177	100.95	3.14	7
R91566 0.25 AA159179 0.18 AA25307 2.25 R59473 0.03 H29290 0.08 AA401434 0.02 AA401434 0.02 AA401434 0.01 R49439 0.01 R49439 0.01 AA460274 0.01 AA460274 0.01 AA425158 0.01 AA425158 0.01 AA425158 0.01 AA425158 0.01 AA701996 0.07 AA701996 0.07 AA701996 0.07 AA701996 0.01 AA701996 0.01 AA701996 0.01 AA701996 0.01 AA701996 0.01 T57803 1.09 AA49045 0.031	37.53	600	52.03	51.22	7
AA159179 0.18 AA425307 2.25 R59473 0.03 H29290 0.08 AA401434 0.02 AA401434 0.02 AA401439 0.01 R49439 0.01 R49439 0.01 AA460274 0.01 AA460274 0.01 AA425158 0.01 AA425158 0.01 AA425158 0.01 AA425158 0.01 AA425158 0.01 AA701996 0.07 AA701996 0.07 AA701996 0.07 AA701996 0.01 AA701996 0.01 AA701996 0.01 AA701996 0.01 T57803 1.09 AA49045 0.031		5	06.63	3.18	2
AA425307 2.25 R59473 0.03 H29290 0.08 AA401434 0.02 AA401434 0.02 AA403529 0.01 R49439 0.21 R90934 0.21 R90934 0.21 AA443624 0.01 AA443624 0.01 AA425158 0.01 AA425158 0.01 AA701996 0.02 R15813 0.01 AA701996 0.02 AA701996 0.01	40.04	4 4	03.87	3.37	8
R59473 0.03 H29290 AA401434 0.02 AA401434 0.02 AA405052 0.26 AA099369 0.01 R49439 0.21 R90934 0.01 AA460274 0.01 AA460274 0.01 AA443624 0.01 H50655 0.63 AA425158 0.01 AA425158 0.01 AA701996 0.02 R15813 0.01 AA701996 0.02 AA701996 0.01 AA701996 0.01 AA701996 0.01 AA701996 0.01 AA701996 0.01 AA701996 0.01 T57803 1.09 AA49045 0.31	32.64	7.07	30.08	3.	
H29290 0.08 AA401434 0.02 AA401434 0.02 AA409369 0.01 R49439 0.21 R90834 0.21 R90834 0.21 H506274 0.01 AA443624 0.01 H50655 0.53 AA425158 0.01 N47445 0.02 R40018 0.07 AA701996 0.02 R15813 0.01 W94486 0.01 H19826 1.00 H96847 0.01 T57803 1.09	32.46	1.79	66.50		• (
AA401434 0.02 AA485652 AA485652 0.26 R49439 0.01 R49439 0.21 R80834 0.21 R49439 0.01 AA460274 0.01 AA443624 0.01 H50655 0.53 AA425158 0.01 N47445 0.02 R40018 0.07 AA701996 0.02 R15813 0.01 W94486 0.01 W94486 0.01 T57803 1.09 AA49045 0.31	31.22	87.92	0.12	5.61	7
AA485052 AA485052 AA485053 B49439 B15740 AA460274 AA443624 C011 AA443624 C011 AA443624 C011 AA425158 C011 AA701996 AA701996 C012 AA701996 C012 AA701996 AA701996 C012 AA701996 C012 AA701996 C012 AA701996 C012 AA701996 C013 AA49045 C011 AA490945 C011	30.56	2.89	81.20	7.60	7
AA099369 R49439 R49439 0.21 R890834 0.21 R800834 0.01 AA460274 0.01 AA4436274 0.01 H50655 0.63 AA425158 0.01 AA701996 C007 AA701996 C007 AA701996 H19826 1.00 H96847 0.01 AA490945 0.31	30.35	0.77	34.54	55.73	61
AA49399 0.21 R49439 0.21 R49439 0.21 R49634 0.01 AA460274 0.01 AA43624 0.01 H50655 0.63 AA425158 0.01 N4745 0.02 R40018 0.07 AA701996 0.02 R15813 0.01 W94486 0.01 H19826 1.00 H96647 0.01 AA490945 0.31	30.08	1.00	7.97	81.28	7
R80834 0.58 R15740 0.01 AA460274 0.01 AA443624 0.01 H50655 0.53 AA425158 0.01 N47445 0.05 N52767 0.02 R40018 0.07 AA701996 0.02 R15813 0.01 H19826 1.00 H19826 1.00 H96647 0.01 T57803 1.09 AA49045 0.31	27.85	1.51	76.38	5.06	2
R15740 0.01 AA460274 0.01 AA43624 0.01 H50655 0.53 AA425158 0.01 N47445 0.02 R40018 0.07 AA701996 0.02 R15813 0.01 W94486 0.01 H19826 1.00 H19826 1.00 H96647 0.01 AA490945 0.31	27.63	72.37	9.84	0.70	2
R15740 AA460274 0.01 AA443624 0.01 H50655 0.63 AA425158 0.01 N47445 0.05 R40018 0.07 AA701996 0.02 R15813 0.01 AH19826 1.00 H96647 0.01 AA490945 0.031	27.34	0.74	70.25	11.03	2
AA4902/4 AA443624	27 13	1.21	69.04	11.15	7
AA42514 0.01 AA425158 0.01 N47445 0.05 N47445 0.05 R40018 0.07 AA701996 0.01 AA701996 0.01 W94486 0.01 H19826 1.00 H96647 0.01 T57803 1.09 AA49045 0.31	27.04	8.33	1.36	71.43	2
AA425158 0.01 N4745 0.55 N52767 0.02 R40018 0.07 AA701996 0.02 R15813 0.01 W94486 0.01 H19826 1.00 H96647 0.01 T57803 1.09 AA49045 0.31	26.99	0.69	27.83	52.45	2
AA420130  W47445  N52767  0.05  R40018  0.07  AA701936  0.02  R15813  W94486  0.01  H19826  1.00  H96647  C01  AA49045  0.31	26.26	5.77	72.00	<b>6</b> .	7
N4745 0.55 N52767 0.02 R40018 0.07 AA701996 0.02 R15813 0.01 W94486 0.01 H19826 1.00 H19826 1.00 A449045 0.31	25.90	2.28	11.75	63.66	2
NS2767 0.02 R40018 0.07 AA070437 0.01 AA701996 0.02 R15813 0.01 W94486 0.01 H19826 1.00 H19826 1.00 H36847 0.01 T57803 1.09 AA49045 0.31	25.65	0.59	66.04	10.31	2
AA701996 0.01 AA701996 0.02 R15813 0.01 W94486 0.01 H19826 1.00 H96647 0.01 T57803 1.09 AA49045 0.31	25.55	2.05	68.60	6.05	7
AA701996 0.02 R15813 0.01 W94486 0.01 H19826 1.00 H35647 0.01 T57803 1.09 AA49045 0.31	25.21	2.29	67.88	5.46	2
AA/01990 C.02 R15813 C.01 W94486 C.01 H19826 1.00 H95647 C.01 T57803 1.09 AA490945 C.31	25.17	3.88	71.30	0.34	7
M3647 0.01 H36647 0.01 T57803 1.09 AA490945 0.31	25.10	2.87	64.40	8.05	7
H19826 1.00 H19647 0.01 T57803 1.09 AA490945 0.31	24.95	2.32	41.17	31.35	71
H96647 0.01 T57803 1.09 AA490945 0.31	24.86	2.27	26.82	45.50	7
H96647 0.03 T57803 1.09 AA490945 0.31	03.450	1 76	60.48	11.25	2
T57803 1.09 AA490945 0.31	24.30	9 7 0	46.00	62.76	7
AA490945 0.31	24.25	400	28.65	43.13	2
	74.24	20 1	300	70 07	, ,
	23.68	0.87	21.91	40.43	4 (
AA490044 0.28	23.31	0.87	<b>4</b> .5	4.51	7

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

7	2	7	۲3	2	2	7	7	7	7	7	7	2	2	2	2	7	7	7	7	2	7	7	2	7	2	7	7	7	7	7	2	2	7	7	
25.34	0.73	53.51	54.20	23.77	3.87	3.43	17.48	3.78	28.27	20.17	50.17	45.41	25.26	49.74	46.16	1.08	6.54	6.22	5.54	16.22	8.77	1.27	12.21	¥.	7.22	5.65	28.36	25.15	3.27	0.57	15.13	18.34	3.12	6.87	
42.67	61.72	9.53	10.05	1.50	58.79	57.89	42.76	0.20	30.54	38.72	7.58	10.31	30.81	1.00	1.93	14.31	47.06	45.91	45.44	34.15	40.61	3.64	37.48	1.51	39.78	40.90	16.52	21.09	41.42	41.20	28.43	26.10	38.63	0.27	
1.00	3.92	2.34	0.88	39.83	0.72	0.93	0.93	56.76	1.29	1.00	1.10	2.79	2.35	7.38	69.6	41.41	2.02	1.21	1.87	1.97	2.37	46.78	1.87	14.65	2.86	2.84	2.63	0.69	1.59	3.92	1.53	0.42	2:32	38.64	
23.00	22.12	21.79	21.71	21.70	21.13	20.75	20.39	20.24	20.04	19.97	19.61	19.50	19.47	19.37	19.26	18.93	18.21	17.78	17.62	17.44	17.25	17.22	17.19	16.78	16.62	16.47	15.84	15.64	15.42	15.23	15.03	14.95	14.69	14.60	16
0.23	0.30	8	3.65	0.14	0.51	0.95	6.19	1.02	3.94	0.20	3.26	0.20	3.30	0.19	0.19	21.01	0.74	3.45	0.54	0.17	4.0	8.76	97:	0.11	0.29	1.14	2.94	5.98	2.60	0.95	3.91	3.94	1.38	0.54	
0.01	0.01	0.02	71.0	500	0 02	0.05	80	0.05	0.20	0.01	0.17	0.0	0.17	0.01	0.01	1.11	2	0.19	0.03	0.0	0.03	0.51	90.0	0.01	0.02	0.02	0.19	0.38	0.17	90.0	0.26	0.26	0.09	9.0	
170198	H88143	AA181007	A4401261	H40072	100111	NZARAB	AA255954	R43910	R95684	AA461508	AA520979	R86970	H28091	AA496022	AA456628	T65736	AA620747	AA487505	R60020	AA001376	AA489662	AA857809	AA857163	AA481481	N26559	N51297	AA504838	AA417713	AA206311	N73083	A463445	H18932	R49592	AA035745	
81050	220655	591055	924270	077470 7KB43	31187	267085	686552																											364865	
510	220	8 6	3 6	3 5	77	<b>1</b>	3 2	200	£ 2	529	230	8	532	533	700	\$2	83	237	838	830	540	3	242	543	3	545	8,48	3	25	548	350	551	552	553	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithellal Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

7	8	ď	2	2	71	c	<b>4</b> (	<b>v</b> (	7	7	<b>7</b>	7	7	7	<b>7</b> . (	~	7	7	7	2	2	2	7	7	7	7	7	7	8	7	7	7	7	7	7	
2.07	23.25	2.72	3.04	4.61	3.01	(F 4)	0.00	90.12	12.18	18.27	29.19	26.50	7.37	25.70	28.39	223	5.10	4.25	0.58	7.02	26.39	20.98	11.09	1.00	4.42	20.10	3.37	16.24	28.00	3.44	14.01	28.79	3.60	3.82	3.43	
38.09	18.93	36.42	37.99	35.90	2.96	200	36.28	<del>.</del> .	26.44	21.58	8.82	<del>2</del> .	97.0	12.28	8.17	29.01	31.49	31.19	14.55	29.00	0.39	14.19	23.87	29.43	2.25	14.95	30.06	17.13	4.46	30.95	20.54	0.92	28.84	27.04	28.86	
3.43	1.00	3.59	5.58	90: 1		1 100	0.88	2.99	2.42	0.64	2.08	10.97	30.80	0.65	1.86	7.14	1.33	2.47	22.57	1.18	10.40	1.96	2.11	6.29	, 29.85	1.00	2.42	2.08	2.89	0.81	0.58	4.71	1.92	2.83	1.36	
14.53	14.40	14.24	14.20	07:5	5.15	14.13	14.09	13.89	13.68	13.50	13.36	13.00	12.97	12.88	12.81	12.79	12.64	12.63	12.56	12.41	12.39	12.38	12.38	12.24	12.17	12.02	11.95	11.82	11.78	11.74	11.71	11.47	11.45	11.23	11.22	17
5.51	41.0	<u> </u>	<b>1</b> (	7,47	0.77	12.46	2.19	4.17	1.31	3.95	8.41	0.08	0.36	0.13	8.32	0.71	1.99	0.33	0.13	4.99	44.	0.12	3.72	0.12	18.67	1.69	0.52	23.53	0.12	0.12	0.18	0.12	0.70	7.46	2.42	
0.38	5	5 6	50.0	0.1/	0.05	0.88	0.16	0.30	0.10	0.29	0.63	0.0	0.03	0.01	0.65	90.0	0.16	0.03	10.0	0.40	0.76	10.0	0.30	0.01	1,53	0.14	90.0	8.	0.0	0.0	000	0.0	90.0	99.0	0.22	
AA863125	0.00000	AA434388	A8453200	R40106	AA010224	N53380	AA699601	W90705	R37738	PA427719	H15549	N52193	AA035730	AA017379	AA911236	R43915	AA070435	H11051	0 A 4 5 7 6 1 8	AA437212	A4620697	AA022498	H10713	157927	AA464617	R92201	AA398321	AA477227	W51795	AA456143	AA161188	W72881	H18015	AA478952	AA609300	
1455603	03001	0980//	4508/	30082	430237	284001	436402	418004	26806	774004	68484	284288	364844	361122	1526789				705446	757428	1049185	364352	47264	71671	812977	195845	79797	739450	324404	706367	501157	345262	50982	753984	1031561	
KKA	Ş :	555	92 92	221	228	<b>22</b> 8	999	561	567	, E	3 3	¥ \$	35.	28.	568	5,60	8 2	5 6	; £	7 6	275	7 2	37.5	215	87.8	573	8 8	, 4 , 4 , 4	5 6	700 784	200	, 85 58 58 58 58 58 58 58 58 58 58 58 58 5	586	587	588	;

Table 2-10; Prostate Cancer Cell Lines vs, Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

2	7	2	7	7	7	7	7	7	7	2	7	8	7	7	5	2	64	7	5	7	7	7	7	2	7	7	2	73	7	7	2	7	7	7	
16.34	7.83	27.27	27.95	13.80	<b>6</b> .00	4.84	16.62	10.20	3.26	20.10	8.36	4.31	21.77	4.40	4.66	2.78	6.78	4:17	15.60	3.50	23.76	15.64	0.48	1.65	1.83	17.42	24.42	1.12	16.30	1.00	15.31	24.03	10.88	1.81	
16.85	22.78	4.45	1.00	17.96	26.77	25.57	15.37	20.21	27.39	10.22	1.00	25.25	8.99	25.88	1.37	23.39	21.63	24.49	14.48	24.58	6.31	14.23	7.21	17.15	25.02	11.85	6.06	20.18	12.15	24.20	13.69	0.98	17.62	19.71	
0.33	2.56	1.10	3.77	0.92	1.90	2.21	0.51	2.09	1.79	2.05	22.91	2.40	0.74	1.14	25.10	4.91	2.67	2.40	0.88	2.87	0.86	0.77	22.87	11.51	3.45	0.83	2.67	8.61	1.40	4.50	0.67	4.65	1.10	8.07	
11.17	11.06	10.94	10.91	10.90	10.89	10.87	10.84	10.83	10.82	10.79	10.76	10.65	10.50	10.47	10.38	10.36	10.36	10.35	10.32	10.32	10.31	10.21	10.19	10.11	10.10	10.01	10.05	9.97	9.95	06'6	9.89	9.83	9.87	9.86	18
9.3	0.95	1.32	0.11	3.90	0.36	1.12	6.12	0.31	3.13	9.32	0.11	96.7	8.09	7.51	18.95	12.67	4.0	0.21	0.87	4.59	4.21	5.55	9.47	0.12	8.21	2.53	16.05	2.42	2.29	0.10	4.94	45.45	3.82	0.09	
0.03	60:0	0.12	0.01	0.36	0.03	0.10	0.57	0.03	0.29	0.86	0.01	0.75	0.77	0.72	1.63	1.22	9.0	0.02	0.08	0.45	0.41	0.54	0.93	Q 201	0.81	0.25	1.60	0.24	0.23	0.01	0.50	4.60	0.39	0.01	
AA160670	AA427737	AA454222	N31808	H48070	AA453787	R66438	AA410434	N74131	N66900	N30751	AA680136	R39765	AA489073	AA463248	AA702803	AA421171	N93721	AA418852	AA281784	AA099148	H54451	N49389	AA412049	AA158035	AA775223	R94504	AA478036	AA626167	AA564007	AA430042	AA292226	R25377	N21633	R55220	
592728	770854	795526	271483	193333	813738	41869	753428	298417	287190	257955	433155	26164	824802	797055	448036	731095	307138	768043	712401	489509	203184	277621	729953	590640	868838	197651	740457	745248	855391	781110	725877	133136	266135	154720	
589	290	591	592	593	594	595	596	597	869	299	8	6	602	603	90	80	909	607	809	609	610	611	612	613	419	615	616	617	618	619	620	621	622	623	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

· • • • • • • • • • • • • • • • • • • •		!						
J 1 W 1		0.54	6.27	9.80	2.04	22.72	4.63	
1 20 14	۹.	<b>t</b> 6	, c	9.79	1.50	24.64	3.25	
· w /		0.20	2 6	27.0	1.35	22.17	5.79	
<b></b>		0.02	17.0	17.6	Ş	9.16	18.56	
	305 AA620528	0.62	5.97	80.0	. ·	2, 22	4.67	
	725 AA447986	0.01	0.10	9.67	S	1 2	28	
-	512116 AA133590	0.64	6.12	9.62	0.66	F.1.2	9 6	
		0.48	4.60	9.57	0.81	24.53	<b>3</b> 7.7	
•		20.0	0.16	9.56	1.61	17.50	9.58	-
		20.0	72.0	9.53	2.05	15.63	10.89	
_		9.0		9.51	24.42	3.74	0.37	
• •		0.70	2 5	0.00	23.58	3,83	1.89	
535 375		1.37	12.98	00.6	1 16	11.85	15.44	
<b>536</b> 530	013 AA219060	0.15	1.40	υ. Α	2 .	8 8	16.97	
	495 W15659	0.01	0.09	9.47	10.45	3. 3	10.01	
		0.01	0.09	9.40	8.70	<u>\$</u>	0.40	
	959 AA136664	0.53	5.02	9:39	1.84	5.32	00.rz	
		0.41	3.84	9.37	0.66	23.58	3.86	
_	40550 AA600006	0.01	0.08	9.36	8.59	1.12	18.37	
		0 03	0.32	9.36	4.85	21.39	1.83	
	•	100	0.06	9.34	11.43	1.54	15.08	
	•		0.22	9.31	0.97	12.53	14.44	
		6 6	8	9.30	4.26	1.86	21.78	
		5 6	8 8	62.5	3.14	1.00	23.74	
		5 6	, w	62.6	1.35	16.72	9.79	
*	791 AA455242	0.0	S. 6	92.6	2.30	7.70	17.71	
		<b>*</b> C	) i	85 6	2.37	19.73	5.45	
-		2 6	5 6	0 13	0.81	13.04	13.51	
_		0.27	<b>4</b> 5	3 60	12.50	0.36	14.35	
••	_	50.0	0.20	, i	S C	100	21.12	
•	•	0.0	0.09	) i	36.	10.87	14.67	
653 286	3259 N26515	0.35	3.21	90.5	5	5 6	- N	
_	527541 AA192527	0.40	3.62	9.05	- - - - -	20.24	0 to	
		1.17	10.62	9.04	2.44	15.22	9.50 04.00	
		021	11.71	9.03	12.54	13.45	1.10	
		0.48	4.35	8.99	1.42	21.38	4.17	
	391033 AA130340	80.	17.81	8.98	10.05	0.30	16.60	
658 858	1241 AA620463	08:-	2	, ,				

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

7	0	2	7	8	8	2	2	7	2	2	7	8	7	8	8	7	74	2	2	7	2	7	7	2	2	7	7	2	2	7	7	2	2	7	
5.40	11.94	8.45	14.43	8.30	18.10	0.61	3.16	7.50	3.36	14.35	18.86	17.99	19.50	11.27	12.36	17.21	2.28	16.24	19.27	2.96	13.09	11.26	0:50	18.07	3.68	9.48	4.38	17.47	5.36	13.46	18.54	8.40	8.68	3.15	
20.84	2.96	17.16	2.08	17.44	8.6 29.	21.88	21.71	16.58	21.85	1.00	0.52	6.50	3.76	12.46	11.51	6.16	18.57	6.27	0.41	13.18	1.67	11,14	19.58	0.56	20.61	13.54	17.84	2.94	18.34	8.34	4.66	14.68	13.94	18.08	
9.65	11.72	0.92	9.92	0.36	<u>4</u> .	3.59	1.16	1.87	0.60	10.33	6.22	0.78	1.96	1.45	1.30	1.81	4.16	2.41	5.22	8.74	9.98	2.23	4.39	5.83	0.11	1.22	1.89	3.70	0.38	2.25	0.81	0.92	1.06	2.43	
8.96	8.87	8.84	8.81	8.70	8.69	8.69	8.68	8.65	8.60	8.56	8.53	8.42	8.41	8.39	8.39	8.39	8.34	8.31	8.30	8.29	8.25	8.21	8.16	8.15	8.13	8.08	8.07	8.04	8.03	8.02	8.00	8.00	7.89	7.89	20
0.21	5.97	3.32	2.16	39.21	0.80	4.61	9.68	34.28	5.05	0.09	5.11	6.98	11.80	9.51	3.36	0.08	0.08	13.45	10.75	1.09	0.15	1.49	0.16	0.14	21.79	6.11	. 16.15	0.09	0.22	0.08	5.71	8.81	2.29	15.64	
0.05	0.67	0.38	0.25	4.51	0.09	0.53	1.12	3.86	0.59	0.01	0.60	0.83	1.40	1.13	0.40	0.01	0.0	1.62	1.30	0.13	0.02	0.18	0.05	0.02	2.66	0.76	200	0.01	0.03	0.0	0.71	1.10	0.29	1.98	
AA416970	W68220	H41203	H16772	AA629999	AA778646	W72671	AA609471	AA775863	R40373	AA496998	AA682321	N80784	T87515	W47364	AA504265	AA036974	AA132226	N74617	AA488324	R61372	N50806	AA058576	R47893	AA131239	AA447504	AA689427	AA701351	AA480961	AA457576	H94944	AA487674	H51765	AA284172	W90381	
730036	342640	175533	50227	884511	1048995	345761	743589	878544	28927	897546	462926	300615	115408	324717	825416	484535	566887	296429	842968	37980	280882	380884	153355	503581	784285	433253	435488	796117	838776	230261	841620	193892	701625	417920	
628	99	961	99	663	189	555	<b>§</b>	299	999	699	670	671	672	678	674	675	676	22	878	679	680	189	682	683	489	685	989	687	688	689	069	691	692	693	

ulated 3 fold or higher In 3 of 3, or 5 fold or higher In at least Table 2-10: Prostate

81	2	7	8	2	2	2	. ~						• •		١ ،	<b>4</b> (	<b>v</b> (	<b>,</b>	7 (	N (	7 (	7	<b>C4</b>	2	7	7	7	2	2	2	8	2	7	7	2	
2.87	3.76	5.01	18.64	0.99	15.20	90	0.28		37.48	5.00	7 6	t 6	3 8	3.16	5.00 9.00	9.32	97.61	2.32	15.99	9.88	15.83	1.19	9.18	. 09.4	12.04	9.29	15.98	3.75	8.02	15.57	14.40	10.67	1.23	10.22	12.41	
	ะ	9.0	18.	ö	10.																										•				·	
16.63	1.50	15.21	3.07	4.09	7.43	1 2	26.71	8.71	14.11	3.07	11.89	14.72	80.9	16.52	16.02	12.70	5.74	17.38	1.00	12.36	5.39	18.13	0.92	15.92	8.97	11.96	4.20	16.45	13.66	4.08	9.0	0.08	16.05	9.92	2.06	i
4.16	18.23	2.27	1 62	101	2 6	7.0	0.97	 81.1	7.96	1.69	5.09	1.41	12.36	2.69	1.23	0.83	1.27	3.02	5.64	0.35	1.25	3.13	12.26	1,82	1.30	0.99	1.89	1.85	9.3	2.29	6.83	11.07	4.52	1.57	 	34.7
7.89	7.83	7 83		0 7	1.14	7.73	7.73	7.70	7.69	7.67	79'2	7.66	7.65	7.64	7.64	7.61	7.60	7.57	7.54	7.53	7.49	7.48	7.46	7,45	7.44	7.41	7.36	7.35	7.34	7.31	7.29	7.27	7.27	4 6	1.64	(7.73
8.53	7 30	6. r	07.6	21.45	1.06	0.14	1.86	2.05	90.0	22.51	0.53	0.25	<del>.</del> 8.	23.26	18.06	0.08	5.46	12.06	0.08	7.28	1.18	5.21	0.08	2.44	4.	0.31	8.77	4	. 6	, a	51.52 74.84	46.41		3.90	0.35	2.20
ğ	3 3	<b>5</b> .	0.66	2.76	0.14	0.02	0.24	0.27	0.0	2.93	0.07	0.03	0.13	3.04	2.36	0.01	0.72	1.59	0.01	0.97	0.16	0.20	200		6.0	800	1 10		2 4	5 +	• •		0.7 6	0.52	0.05	0.30
	N62132	AA826324	R50775	W74362	H14949	AA156795	AA453437	H57082	AA463982	AA700739	H10231	H10068	AA780365	AAURRRAG	W72666	T95804	AA291773	B59197	00000000	P52455	4641415	7700000	AA/00/40	CTOCOTA	AA/028/3	K164/3	A A E 3 4 3 6 4	•	•		•	166816	AA706968	R42946	N22766	R43308
	287598	1420830	38833	345090	48330	502444	788217	204644	810661	435351	46927	46740	1035664	581971	345743	120695	725340	41789	448033	30833	988160	9000	200000	001110	44/16/	C#7071	90000	07000	428218	278729	614260	6631/	451907	31239	266631	22755
	<del>2</del>	695	969	697	698	669	92	707	5	5	2 2	5 5	3 8	3 5	5 5	3 5	B 5	2 ;	= ;	77	2 ;	<b>*</b> ;	55	3,18	71	78	2 2	3	22	22	723	24	725	728	727	728

Table 2-10: Prostate Cancar Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

8	7	7	7	7	7	7	7	7	7	7	7	7	7	7	7	7	7	2	7	74	71	7	7	7	7	7	7	2	2	7	7	2	7	7	
6.82	6.11	6.43	11.27	15.36	16.05	14.81	10.38	11.80	7.84	7.62	5.32	8.84	0.77	6.81	0.27	4.28	11.87	14.49	3.14	9.27	10.38	15.45	1.16	8.65	12.53	8.19	13.46	8.34	11.61	3.78	5.09	6.48	15.52	3.19	
13.06	12.65	13.46	7.61	3.28	3.21	4.52	8.83	8.75	0.03	11.28	2.34	11.35	15.66	12.50	17.83	14.80	7.50	5.67	16.93	9.22	8.38	1.38	5.14	8.72	1.10	1.51	2.38	8.48	7.15	2.82	13.88	12.71	3.80	16.50	
1.81	2.92	1.78	27.2	2.94	2.29	2.42	2.25	0.87	13.51	2.47	13.68	1.14	4.89	2.00	3.0	1.98	1.68	0.85	0.86	2.37	2.09	3.97	14.50	2.34	7.04	10.95	4.74	2.73	1.77	13.90	1.52	1.22	1.08	0.68	
7.23	7.23	7.22	7.20	7.19	7.18	7.18	7.17	7.14	7.13	7.12	7.11	7.11	7.11	7.11	7.05	7.02	7.02	7.00	6.98	6.95	6.95	6.94	6.93	6.90	68.9	6.88	6.86	6.85	6.84	6.83	6.83	6.80	6.80	6.79	23
1.17	8.99	0.31	4.0	11.27	8.58	0.24	15.96	0.11	105.24	14.44	0.62	1.51	3.14	95.95	0.26	2.95	2.01	0.28	1.75	0.68	36.26	0.05	0.07	3.78	90.0	2.74	23.86	4.75	0.83	3.35	0.25	2.79	4.70	5.63	
0.16	124	90.0	9.6	1.57	1.19	0.03	223	0.02	14.71	2.03	0.09	0.21	4.0	13.50	30	0.42	0.29	9.0	0.25	0.10	5.22	0.01	0.01	0.55	0.01	0.40	3.48	0.69	0.12	0.49	9.0	0.41	69.0	0.83	
H19234	AA135809	AA011182	AA455521	AA431773	AA131909	AA128005	AA504201	AA677655	AA894557	AA455237	AA449118	R96520	AA455087	N66158	AA029722	AA598849	N53940	W72556	AA417288	168878	AA485453	AA488741	T58146	AA886739	AA669222	AA452278	T83880	AA129135	A1025259	AA935560	AA398406	H56903	R23246	AA455940	
50904	565734	369722	809828	782503	504308	501868	825282	897271	1416782	814779	785845	199644	812294	278531	366778	898333	247050	345330	731139	82215	811067	841221	79254	1500162	856585	787938	79781	586650	1641367	1556526	726791	204580	131094	813275	
729	730	734	732	733	75	735	238	737	7.38	739	740	741	742	743	74.	746	746	747	748	749	750	751	752	753	754	755	220	157	758	759	200	761	762	763	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

	2	2	7	,		1	2	74	2	2	7	7	ч	7	73	71	7	7	7	7	7	73	2	8	2	7	7	7	7	7	7	7	7	2	7	7		
	1.76	0.98	883	,	3.16	55.7	2.21	13.94	14.72	4.64	4:14	4.12	8.09	1.85	9.54	45.	4.81	1.28	9.68	6.74	13.83	5.55	6.04	3.84	2.39	12.67	7.59	8.78	6.08	9.70	1.28	12.22	4.18	8.37	3.99	12.04		
	15.10	70	64.	1.82	16.41	10.73	14.81	4.25	3.97	14.02	13.65	1.82	0.58	10.05	9.16	12.93	13.90	13.84	7.59	11.33	4.25	11 59	2.65	13.27	3.01	184	9.29	9.13	11.14	6.70	12.25	6.06	1.59	7.88	13.18	5.53		
	37 6	0.4.0	12.08	9.56	0.64	2.08	3.11	1.87		5. to	c	17.51	14.01	1 07	1.16	- c	4 02	7 79 7	10.4	¥ ;	1.4	Ç	2.35	10.78	\$ 70°	5.6	08.4	5. 5. 8. 5.	2.05	2 85	69	06.0	13.36	2.87	 	\$ <del>\$</del>		
	1	6.77	6.77	6.74	6.73	671	0.71	0.7.0	6.03	6.68	6.67	6.67	6.65	49.9	5.62 	5.62	85.58 8	6.58	6.58	6.57	6.51	6.51	6.50	6.49	6.49	6.48	6.47	6.46	5.45	0.42	4. 4.	- c	0.0	9.3/	6.37	6.37	6.57	<b>c7</b>
		7.09	1.65	10.08	88.0	3 1	4.7	3.87	421.08	3.42	3.96	3.02	17.94	7.75	0.02	63.60	2.64	1.48	1.57	0.12	0.24	5.27	6.55	8.71	10.14	2.76	0.07	0.12	27.63	0.45	6.74	0.21	99.1	9.86	3.76	0.11	4. 4.	
		1.05	0.24		3 !	1.4/	1.11	0.58	62.96	0.51	0.59	0.45	2.70	1.17	0.01	9.61	0.40	0.22	0.24	0.02	0.04	0.81	1.01	1.3	1.58	0.43	0.01	0.02	<b>6.3</b>	0.07	1.05	0.03	0.26	- - - - -	0.59	0.02	0.71	
•		M48707	2000	700 MONAY	AA454597	N47008	AA398949	H05085	AA857413	R53446	W37447	AA883391	AA165678	AA424790	AA448400	155728	AA142869	AA029997	H09997	AA039851	T51617	N79745	R43328	AA953357	AA598796	R32406	AA129677	AA428186	AA626788	AA452542	AA448664	H15910	R89846	N25936	N39611	AA485351	AA405533	
		700070	102017	COCAOR	811582				1475028	40038	321900											300276	32134	1573251	898088	135688	565493	773564	87778	788488	786078	159462	194717	268711	277112	810996	772410	
			\$		992	767		_		3 5					77.6								3 5				٠.								796	797	798	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

~ (	7	2	7	7	2	2	7	2	2	7	7	7	2	7	2	2	2	7	2	7	2	7	7	2	7	7	7	2	7	7	7	2	2	7	
6.13	11.23	1.13	11.81	9.86	6.32	8.17	7.26	8.95	7.88	10.08	13.49	5.30	1.69	5.48	9.06	99.6	3.89	6.77	2.74	6.60	7.78	10.29	9.56	9.04	3.85	1.49	7.48	7.07	11.91	5.03	1.00	12.34	12.62	4.79	
12.20	6.38	4.43	4.38	8.03	10.78	8.05	10.13	2.46	8.82	6.21	5.01	0.27	3.53	12.34	8.92	7.60	12.42	0.76	3.72	10.69	2.18	7.55	2.69	7.76	13.05	12.18	9.73	8.86	3.64	11.73	3.46	4.07	0.62	11.45	
0.76	1.48	13.51	2.86	1.06	1.79	2.64	1.45	7.40	2.11	2.51	0.27	13.18	13.52	0.87	0.68	1.35	2.28	11.05	12.11	1.20	8.48	0.57	6.15	1.59	1.47	4.68	1.12	2.39	2.76	1.52	13.72	1.70	4.85	1.83	
6.36	6.36	6.36	6.35	6.31	5	62.9	628	6.27	6.27	6.27	6.25	6.25	6.24	6.23	622	6.20	6.20	6.20	6 19	6 16	 	6.13	6.13	6.13	6.12	6.12	6.11	6.11	6.10	6.09	90.9	6.03	6.03	6.02	24
1.50	6.46	3.38	26,80	4.87	5 5	5. FR 3	i c	13.7	ζ. ·	0.15	2.29	13 80	86.4	4 11	<b>.</b> .	5 01	90,	800	8 8	5 5	3 5	15.5 19.4	12.43	2 89	10.01	0.15	2.40	16,02	207.98	26.54	900	0.70	5.33	16.34	
0.24	101	0.53	4 22			C7.50	2 6	C. 53	77.0	67.0	5 6	3 6	. C	2 2	3 6	77.7	5 .	<b>.</b>	5 5	? ?	6.00	1 C	<b>6</b> .00	270		200	0.39	2.62	80.75	4.36	100	0 12	1 88 5 C	271	<b>.</b>
R93068	A A 455773	AA777655	44664	188171	AA459244	AA042812	W85878	N63988	N77828	H54659	10788	NC4009	AA430367	A 6 4 7 6 4 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9	0/810/90	KC8338	000771	AA65U3/3	W31389	AA//84	9/89CN	AA865924	AA030104	AA430744	A A & O & 7 5 7	DYRAKE	R92188	A A 430052	AA410517	A & 4 5 9 1 0 5	DAAR17	44774547	AA165313	AA453441	
196824	047344	112410		and the second						203179																								788225	
700	2 6	3 3	<b>a</b> 8	305	8	\$	8	8	801	808	3	810	2	719	613	4 4	315	818	817	25 26	8 9	820	821	822	3	Š	3 2	27	3 6	200	3 8	3 5	ē :	מ בנ	3

Table 2-10: Prosta

at least	<u>-</u>	
state Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.	ппаниничинаничиничиничиничин	7
her in 3 of 3, or t	4.18 2.56 2.56 7.27 7.79 4.51 11.52 6.52 8.76 8.76 8.76 8.76 8.76 8.76 8.76 8.76	5.77
d 3 fold or hig sell lines.	9.92 10.65 9.75 7.62 8.57 11.00 3.88 9.04 11.46 0.30 7.34 8.18 11.44 8.58 6.81 12.13 2.32 12.13 12.13 12.13 10.89 10.89 10.89 10.89 10.45 11.11	9.02
mal Prostatic Epithelial Cell Line: Genes upregulated 3 fold 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.	2.14 2.89 4.63 4.63 2.44 1.57 1.157 1.141 1.15 1.15 1.15 1.15 1.15 1.15 1.1	2.42
Cell Line: Gen er in at least 1	6.00 6.00 6.00 6.00 6.00 6.00 6.00 6.00	5.74 5.73 25
ic Epithelial ( fold or high	0.39 0.39 8.34 8.89 8.34 2.99 7.54 0.49 0.55 0.05 0.30 1.67 4.03 0.14 0.06 0.22 0.25 0.25 0.39 0.41 0.06 0.41 0.06 0.25	3.37
mal Prostati 2 of 3 or 10	0.07 1.29 0.09 1.39 0.08 0.09 0.12 0.01 0.04 0.07 0.03 0.03 0.03 0.03 0.03 0.03 0.03	1.07
nes vs. Nor	R59369 AA458646 AA4055840 AA465353 AA504202 N39233 H15696 H08210 AA879423 AA101173 N52868 N36008 AA429474 AA429474 AA429474 AA708608 AA65090 N69068 AA65090 N69068 AA65090 N69068 AA65090 N69068 AA65090 N69068 AA10619 AA10619 AA10619 AA10619	R50752 H10788
cer Cell Li	38027 813410 813410 814080 825284 276872 45607 1505686 583660 289645 272888 796366 28473367 429376 1333468 488516 134395 246549 773367 429376 1331113 506483 36480 26563 287349 287349 287384	
state Can	8.35 8.35 8.36 8.37 8.41 8.42 8.43 8.44 8.43 8.44 8.43 8.44 8.43 8.43	868

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least

5	137847	0.24	1.35	5.72	0.89	6.27	10.01	7
13013	13/01/		06	5.71	1.62	12.44	3.07	8
27817	D40481	0.03	0.15	5.71	2,14	9.62	5.36	2
24063	R45451	0.28	1.58	5.68	572	9.78	1.55	7
291374	N72274	8.	5.71	5.68	1.00	6.51	9.54	7
282104	N51498	0.51	2.92	5.68	5.83	9.35	1.87	7
503865	AA131707	1.39	7.88	5.68	1.17	12.71	3.16	7
75096	PAGAS	041	2.35	5.68	1.17	8.12	7.73	7
878600	AA775259	1.27	7.19	5.67	1.02	4.30	11.69	7
429517	AA011389	0.0	0.06	5.67	3.76	1.00	12.25	7
39219	B51382	0.14	0.81	5.66	3.85	12.66	97.0	2
33474	AA599560	3.28	18.53	5.65	8.40	2.36	6.19	7
856592	A 4 6 8 9 7 1 8	1 19	6.68	5.64	10.55	1.37	5.00	7
395609	AA757588	0.05	0.25	5.62	0.55	7.77	8.55	7
626022	AA427400	8	0.10	5.62	5.02	0.44	11.39	2
357344	W93500	10.0	90.0	5.62	5.85	8.42	2.59	7
869375	AA679907	900	0.18	5.61	1.75	11.49	3.58	7
950926	AA608730	3.17	17.75	5.60	8.06	1.28	7.48	7
361688	W96197	0.0	0.0	5.60	0.80	10.00	5.99	7
193139	H47315	9.0	0.17	5.59	1.43	9.88	5.46	7
121798	198201	0.72	4.03	5.59	6.89	7.41	2.47	7
89049	AA452909	1.97	11.01	5.58	1.16	11.14	4.42	7
43977	H05563	0.70	3.89	5.55	11.02	1.27	4.36	7
471707	AA035580	99.0	3.65	5.55	6.12	2.85	7.67	7
783987	AA443286	0.01	0.05	5.53	3.23	12.83	0.55	7
878788	AA670408	23.88	132.57	5.53	1.02	7.04	8.52	7
505898	AA907714	0.07	0.41	5.52	2.10	8.01	6.45	7
130858	R22206	0.99	5.42	5.50	2.27	10.54	3.69	7
839081	AA487608	12.19	66.94	5.49	7.57	0.09	8.81	8
30845	R22308	2.02	11.11	5.49	0.87	7.74	7.85	7
241432	H80685	1.63	8.93	5.48	7.11	7.64	1.70	~
79556	H51425	0.01	0.05	5.48	3.34	1.00	12.10	И
97206	R92812	0.98	5.38	5.47	0.67	8.14	7.61	7
825408	AA504262	1.29	7.03	5.47	1.85	10.90	3.66	7
525799	AA074446	0.41	2.26	5.46	4.45	05.1	10.42	8

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

	7	2	2	2	7	7	,		7	7	7	2	2	7	7	7	2	2	7	7	7	2	. ~		<b>y</b> (	7	~	2	7	7	7	7	7	2	7	2	7		
	6.30	7.49	8.44	5,51	8.69	7.71	44 94	19.61	3,53	4.35	8.77	4.15	11.07	8.24	4.21	6.03	7.16	17.7	10.03	60.6	5.20	1 6	7.10	3.27	6.95	10.80	7.50	7.02	2.2	8.43	6.43	7.68	0.55	1.88	3.74	1.61	49.9		
	8.29	6.97	6.79	1 A	200		60.03	3.31	10.54	10.79	5.32	10.96	1.00	6.68	10.44	101	7.63	7.20	3.75	9 4	5.0d	0.6	6.76	8.00	6.68	0.18	2.95	2.97	4.37	5.07	7.32	5.49	10.92	10.57	9	5 6 7	. 6 8	Sa. C	
	1 75	8 8	8 9	97.	2.62	1.61	 	1.10	2.12	401	2.06	÷ 01	50.4	3 7	3.5	<b>1</b> 20 0	- 44	5	7.02	2.14	9.04	1.07	1.93	2.59	2.22	4.81	5.35	57.5	0.36	2.21	8	5.50		07.4	200	30.0b	3.27	2.85	
	5.45	0.40	5.44	4.	5.43	5.42	5.41	5.40	5 40	30	8 00° 10° 10° 10° 10° 10° 10° 10° 10° 10°	0.30	5.38	0.00	5.55	5.33	5.32	5.31	5.31	5.30	5.30	5.29	5.29	5.29	5.28	5.27	5 S	35 3	5.20	5.24	17.0	5.23	5.23	5.22	5.21	5.20	5.20	5.19	27
•	•	0.26	9.05	1.05	0.45	33.51	0.59	90	÷	0.12	2.76	7.13	11.29	9.05	11.61	0.28	0.05	6.28	2.11	14.65	21.66	4.03	0.23	20.12	0.41	4 40	j (	3.60	4.0	0.10	0.58	44.95	15.25	90.0	1.87	0.76	0.46	0.35	
		0.05	0.01	0.19	0.08	6.18	0.11		2 6	0.02	0.51	1.32	2.10	0.01	2.17	0.05	0.01	1.18	0.40	2.76	4.09	0.76	9	181	5 6	0 10	0.85	0.70	0.03	0.03	0.11	8.60	2.82	0.02	0.36	0.15	0.09	0.07	
•		AA428514	H96356	R89287	AA884428	AA706829	0050004	A4000 100	AA969184	173031	H18630	AA458827	AA497034	R44292	AA402965	W72310	R73525	R16676	AA598668	AA775865	AA455652	NOTES	NAZRAR	200104	048//	AA628190	W72679	H59305	AA879474	N20338	AA620614	AA133168	AA169645	AA404290	AA284280	W32272	AA121504	AA446748	ţ
		773170	256720	195786	1466998			146//21	1580342	85561	50839	814350	823566	34462	741790	345077	156473	129664	898258	878550	813997	266402	247089	20000	108395	1055831	345787	204299	1505735	264646	1048685	490753	594517	758360	324665	321386	489881	784126	! •
		\$	908		8 8	8 8	3	<b>606</b>	910	911	912	913	914	915	916	917	918	919	6.00	5	25.0	776	3, 6	476	925	926	927	928	929	830	931	932	933	934	935	938	937	8 6	<b>&gt;</b>

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

,,, ,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	939	845602	AA644334	1.83	9.47	5.18	1.72	6.91	6.90	
11         726668         AA398218         0.01         0.05         5.15         7.12         1.00           12         56688         AA398218         0.01         0.05         5.15         4.21         1.15           13         561824         AA128931         4.32         22.27         5.15         4.21         1.15           14         26324         R.1880         0.04         0.07         5.11         0.72         1.108           2         23536         R.38189         0.04         0.19         5.11         1.62         8.40           4         185274         AA14860         6.40         32.60         5.09         1.14         9.02           6         135573         R71880         0.39         1.36         5.10         1.62         8.40           1         44920         AA77773         0.08         0.43         5.10         1.62         8.40           6         15527         AA77773         0.08         0.43         5.10         1.62         8.40           1         43527         AA14860         6.40         32.60         5.09         1.14         8.23           2         445270         AA48060	3	768146	AA426561	0.10	0.51	5.17	1.23	9.14	5.15	
12         501824         AA128931         4.32         22.27         5.15         4.21         1.15           13         418506         WR4774         0.01         0.06         5.15         8.70         0.80           14         205322         H98997         1.54         0.01         0.06         5.15         1.62         3.54           15         23658         H98997         1.54         0.01         0.07         1.106         0.08         0.35         1.10         0.72         1.106         0.02         0.04         0.14         5.10         1.62         3.54           1         448020         AA747740         0.08         0.43         5.10         1.14         9.02         1.106           1         15372         AA148606         6.40         32.60         5.09         1.14         9.02         8.02         8.02         8.02         8.02         8.02         8.02         8.02         8.02         8.02         8.02         8.02         8.02         8.02         8.02         8.02         8.02         8.02         8.02         8.02         8.02         8.02         8.02         8.02         8.02         8.02         8.02         8.02	2	726658	AA398218	0.01	0.05	5.15	7.12	9.1	7.33	
13         418806         W84774         0.01         0.06         5.15         6.70         0.80           14         265342         H98997         1.94         9.93         5.12         1.82         3.54           15         23568         R78169         0.01         0.07         5.11         1.68         5.30           15         23568         R78169         0.04         0.07         5.11         1.68         5.30           1         44820         AA718605         6.40         0.23         1.98         5.09         1.14         9.02           1         44820         AA718605         6.40         32.60         5.09         1.14         9.02           1         4502         4.07         5.10         1.62         8.40         8.23           1         73247         AA418605         0.04         0.22         5.04         0.82         5.23           2         1.0257         AA1860         0.04         0.22         5.04         0.82         5.23           2         2.041         AA2120         8.07         4.076         5.05         1.54         8.12           2         2.041         A.22         <	845	501834	AA129931	4.32	22.27	5.15	4.21	1.15	10.09	
4         262342         H99997         1:94         993         5:12         1:82         3:54           6         23556         R38163         0.01         0.07         5:11         0.72         11.06           8         123016         AAA83338         0.04         0.19         5:11         1.68         5:30           8         448320         AAA14806         6.40         32.60         5:10         1.14         9:02           9         15525         AA14806         6.40         32.60         5:09         1:14         9:02           1         48920         0.39         1:36         5:09         1:14         9:02           1         5523         4892         0.39         1:14         9:02         1:14         9:02           1         5524         AA4196         0.39         3.45         5:04         0.83         5:29           2         1.6552         N4496         0.59         3.45         5:04         0.83         5:29           2         1.6563         A54063         1.71         8.53         4.97         1:24         8.23           2         2.0652         H6152         0.39         1.	943	415806	W84774	0.01	90.0	5.15	6.70	0.80	7.5	
5         23656         R38169         0.01         0.07         5.11         0.72         11.06           6         1293016         AA683338         0.04         0.19         5.11         1.69         5.30           7         448920         AA777779         0.08         0.43         5.10         1.62         8.40           8         491524         AA748605         6.40         32.60         5.09         1.14         9.02           9         155235         RR1889         0.39         1.96         5.09         1.14         9.02           9         155253         RR1889         0.39         1.96         5.09         1.14         9.02           1         15225         RR42109         AA42120         0.34         0.36         0.35         0.39         1.83         0.31         0.35         0.82         0.85         0.35         0.82         0.85         0.85         0.85         0.85         0.85         0.85         0.85         0.85         0.85         0.85         0.85         0.85         0.85         0.85         0.85         0.85         0.85         0.85         0.81         0.82         0.85         0.85         0.81 <th< td=""><td>3</td><th>263342</th><th>198997</th><td>2</td><td>9.93</td><td>5.12</td><td>1.82</td><td>3.54</td><td>10.01</td><td></td></th<>	3	263342	198997	2	9.93	5.12	1.82	3.54	10.01	
448920         AAAT77779         0.04         0.19         5.11         1.68         5.30           8         491524         AAAT4860S         6.40         32.60         5.09         1.14         9.02           8         491524         AAA4860S         6.40         32.60         5.09         1.14         9.02           9         155573         R71889         0.39         1.96         5.09         1.14         9.02           1         1602209         AA902241         3.26         16.55         5.07         6.08         8.82           2         1602202         AA902241         3.26         1.54         8.12         8.25           2         1602202         AA902202         0.04         0.22         5.04         0.82         8.25           2         24162         AA40800         0.99         3.45         5.02         1.53         8.25           3         243653         AA198         1.71         8.53         4.97         1.25         8.53           4         20602         AA40800         0.39         1.93         4.97         1.25         8.63           5         24162         AA40846         1.21	945	23658	R38169	0.0	0.07	5.11	0.72	11.06	3.56	
448920         AATTTT9         0.08         0.43         5.10         1162         840           8 491524         AAA1860S         6.40         32.60         5.09         1.14         9.02           1 155575         R71689         0.39         1.96         5.09         1.14         9.02           1 162209         AA962541         3.26         1.56         5.03         1.14         9.02           2 1 17324         AA480605         0.04         0.22         5.04         1.54         8.12           2 2 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	848	1293016	AA683338	900	0.19	5.11	1.68	5.30	834	
4 491524         AAA148606         6.40         32.60         5.09         1.14         9.02           1 152575         R71689         0.39         1.96         5.09         2.35         5.29           1 1602209         AA9062541         3.26         1.65         5.07         6.08         8.82           2 1602200         AA9062541         3.26         1.65         5.04         0.82         5.29           2 243653         AA90605         0.04         0.22         5.04         0.82         8.65           3 243653         AA90806         0.04         0.22         5.04         0.82         8.65           4 206052         H61552         0.91         4.54         4.99         0.91         8.28           5 604779         AA148037         1.71         8.53         4.97         1.25         6.10           5 826603         AA604631         6.33         1.93         4.97         1.25         6.05           8 826603         AA418040         0.39         1.93         4.97         1.25         6.08           8 826603         AA478642         1.78         8.24         4.95         0.91         6.03           1 86122         AR3040<	24	448920	AA7777779	90'0	0.43	5.10	1.62	8.40	5.28	
4         155575         R71689         0.39         1.96         5.09         2.35         5.29           1 (50220)         AA902241         3.26         16.55         5.07         6.08         8.82           2 (50220)         AA9022541         3.26         16.55         5.04         0.08         8.02           2 (50220)         AA4026050         0.04         0.22         5.04         0.82         8.65           3 (24365)         AA4026050         0.04         0.22         5.04         0.82         8.65           4 (206052)         H61552         0.91         4.54         4.99         0.91         8.28           5 (5427)         AA148637         1.71         8.53         4.97         7.00         6.10           5 (5427)         AA4148637         1.71         8.53         4.97         7.00         6.10           5 (5427)         AA418600         0.39         1.93         4.97         1.25         6.08           8 (5412)         AA41400         0.39         1.93         4.97         1.93         6.10           1 (6412)         H81612         1.78         8.75         4.97         1.93         6.03           1 (6412) </td <td>28</td> <th>491524</th> <th>AA148505</th> <td>6.40</td> <td>32.60</td> <td>5.09</td> <td>1.14</td> <td>9.02</td> <td>5.12</td> <td></td>	28	491524	AA148505	6.40	32.60	5.09	1.14	9.02	5.12	
6         66.02209         AA421230         3.26         46.55         5.07         6.08         8.82           2         824109         AA420230         8.07         40.76         5.05         1.54         8.12           2         824109         AA480805         0.04         0.22         5.04         0.82         8.65           3         243653         N49899         0.69         3.45         5.02         1.63         5.51           4         206052         H61552         0.91         4.54         4.99         0.91         8.28           5         50427         AA149637         1.71         8.53         4.99         7.00         6.10           5         50427         AA149637         1.71         8.53         4.99         7.00         6.10           5         50420         AA149637         1.71         8.53         4.99         7.00         6.10           6         50440         0.33         1.93         4.97         1.26         6.10         6.10           1         26613         4.54         4.99         7.00         6.10         6.10         6.24         6.26         6.10         6.10         6.10	25	155575	R71689	0.39	1.98	5.09	2.35	5.29	7.63	
2 824109         AA421220         8.07         40.76         5.05         1.54         8.12           2 824109         AA480605         0.04         0.22         5.04         0.82         8.65           3 243653         NA9809         0.69         3.45         5.02         1.63         5.51           4 206052         H61552         0.91         4.54         4.99         0.91         8.28           5 504279         AA149637         1.71         8.53         4.97         1.25         6.10           5 825603         AA504631         5.33         26.53         4.97         1.25         10.25           1 825603         AA504631         5.33         26.53         4.97         1.25         6.10           1 826102         NZ0480         0.01         0.05         4.94         8.10         6.10           1 86132         H35600         0.01         0.05         4.94         8.10         6.08           1 86132         H35600         0.01         0.05         4.94         8.10         6.10           1 14722         R82041         0.06         0.30         4.91         1.93         6.93           1 14722         R82041	88	1602209	AA962541	3.26	16.55	5.07	6.08	8.82	0.32	
2 824109         AA490605         0.04         0.22         5.04         0.82         8.65           3 243653         N49899         0.69         3.45         5.02         1.63         5.51           4 206602         H61522         0.91         4.54         4.99         0.91         8.28           5 50427         AA149637         1.71         8.53         4.97         1.25         6.10           5 50427         AA449637         1.71         8.53         4.97         1.25         6.10           5 50427         AA449637         1.71         8.53         4.97         1.25         6.10           1 64132         H39560         0.01         0.05         4.97         1.29         6.33           1 66132         H39560         0.01         0.05         0.34         4.94         8.10         6.10           1 66132         H39560         0.01         0.05         0.30         4.94         8.10         6.33           1 66132         H39560         0.01         0.05         0.30         4.94         8.10         6.33           1 14725         H86042         0.06         0.30         4.91         4.99         8.14	951	739247	AA421230	8.07	40.76	5.05	1.54	8.12	5.48	
4         2043653         N49899         0.69         3.45         5.02         1.63         5.51           4         206052         Hei552         0.91         4.54         4.99         0.91         8.28           5         504279         AA149637         1.71         8.53         4.99         7.00         6.10           5         504279         AA419637         1.71         8.53         4.97         1.25         6.10           5         504272         AA4504631         5.33         26.53         4.97         1.25         10.25           7         264162         N20460         0.39         1.93         4.97         1.39         6.10           1         26412         H33560         0.01         0.05         4.94         8.10         6.13           1         16613         A44845         1.78         8.75         4.94         8.10         6.14           1         16613         A44854         1.76         8.24         4.94         8.10         6.14           2         16613         A44854         1.76         8.24         8.10         8.14         8.14           2         147825         R82041	952	824109	AA490605	9.0	0.22	5.04	0.82	8.65	2.67	
4         206052         H61552         0.91         4.54         4.99         0.91         8.28           5 504279         AA149637         1.71         8.53         4.99         7.00         6.10           8 256023         AA49637         1.71         8.53         4.97         1.25         1.025           8 25602         AA40631         5.33         26.53         4.97         1.25         1.025           1 264162         N20480         0.39         1.93         4.97         1.93         6.33           1 47824         R44346         1.28         6.34         4.95         0.98         8.07           1 47825         R82041         0.06         0.30         4.91         1.91         6.08           1 47825         R82041         0.06         0.30         4.91         1.91         6.08           2 77545         N56082         0.43         2.11         4.90         0.97         8.14           2 5014         H16743         1.70         8.32         4.90         0.37         7.99           2 5014         H16743         1.70         8.32         4.90         0.37         7.99           4 071         H16743	953	243653	N49899	69.0	3.45	5.02	1.63	5.51	7.92	
5         504279         AA149637         171         8.53         4.99         7.00         6.10           5         825603         AA504631         5.33         26.53         4.97         1.25         10.25           7         264162         N20480         0.39         1.93         4.97         1.29         6.13           8         34234         R44346         1.28         6.34         4.95         0.98         8.07           1         186132         H39560         0.01         0.05         4.94         8.10         5.74           1         186132         H39560         0.01         0.05         4.94         8.10         5.74           1         186132         H39560         0.01         0.05         4.91         1.91         6.08           1         147825         R82041         0.06         0.30         4.91         1.20         8.12           2         277545         N56982         0.43         2.11         4.90         1.20         8.72           3         5014         H16743         1.70         8.32         4.90         1.97         8.45           4         4021         H16743	<b>9</b> 58	206052	H61552	0.91	4.54	4.99	0.91	8.28	5.79	
8 25603         AA504631         6.33         26.53         4.97         1.25         10.25           7 264162         N20480         0.39         1.93         4.97         1.59         6.33           3 4294         R44346         1.28         6.34         4.95         0.98         8.07           1 186132         H39560         0.01         0.05         4.94         8.10         5.74           1 147825         R82041         0.06         0.30         4.91         1.91         6.08           1 147825         R82041         0.06         0.30         4.91         1.91         6.08           2 277545         R82041         0.06         0.30         4.91         1.91         6.08           2 277545         R82041         0.04         0.03         4.90         1.20         8.14           3 50114         H16743         1.70         8.32         4.90         0.37         7.89           4 50241         A16743         1.70         8.32         4.90         0.37         7.89           4 504452         A404569         0.14         4.89         1.97         7.62           4 60750         A60460         0.30         1.48<	955	504279	AA149637	1.71	8.53	4.99	7.00	6.10	1.87	
7         264162         NZO480         0.39         1.93         4.97         1.99         6.33           34294         R44346         1.28         6.34         4.95         0.98         8.07           1 186132         H39560         0.01         0.05         4.94         8.10         5.74           1 784772         AA478542         1.78         8.75         4.92         0.97         8.14           1 147825         R82041         0.06         0.30         4.91         1.91         6.08           2 277545         N56982         0.43         2.11         4.90         1.20         8.72           2 277545         N56982         0.43         2.11         4.90         1.20         8.72           2 277545         N56982         0.03         2.11         4.90         1.20         8.72           3 5014         H16743         1.70         8.32         4.90         0.37         7.89           4 5015         A04045         0.04         0.71         4.89         1.97         8.45           4 40721         R5575         0.30         1.49         4.81         2.09         5.39           4 40727         A04043228	996	825603	AA504631	5.33	26.53	4.97	1.25	10.25	3.42	
3 4294         R44346         1.28         6.34         4.95         0.98         8.07           1 86132         H39560         0.01         0.05         4.94         8.10         5.74           1 78472         AA478542         1.78         8.75         4.92         0.97         8.14           1 147825         R82041         0.06         0.30         4.91         1.91         6.08           2 277545         N56982         0.43         2.11         4.90         1.20         8.12           2 277545         N56982         0.43         2.11         4.90         1.20         8.72           3 5014         H16743         1.70         8.32         4.90         0.37         7.89           1 50415         A4904869         0.14         0.71         4.89         1.87         8.45           4 0721         R55750         0.30         1.49         4.89         1.87         7.62           4 0721         R55750         0.30         1.49         4.89         1.57         7.62           4 0721         R5750         2.16         10.43         4.84         2.45         5.44           2 0550         A 80         2.15	798	264162	N20480	0.39	1.93	4.97	1.99	6.33	6.61	
166132         H39560         0.01         0.05         4.94         8.10         5.74           784772         AA478542         1.78         8.75         4.92         0.97         8.14           1 47825         R82041         0.06         0.30         4.91         1.91         6.08           1 47825         R82041         0.06         0.30         4.91         1.91         6.08           2 502201         AA114106         0.02         0.08         4.90         5.15         7.27           3 50114         H16743         1.70         8.32         4.90         0.37         7.89           4 50415         AA904369         0.14         0.71         4.89         1.87         6.45           4 6071         R55750         0.30         1.49         4.89         1.87         7.82           4 6072         AA035384         7.11         34.71         4.88         1.57         7.82           4 8678         AA054328         5.90         28.69         4.81         2.45         5.47           8 80508         AA64554         0.01         0.05         4.80         7.91         5.50           3 2517         R43571         0.01 <td>958</td> <th>34294</th> <th>R44346</th> <td>1.28</td> <td>6.34</td> <td>4.95</td> <td>0.98</td> <td>8.07</td> <td>5.79</td> <td></td>	958	34294	R44346	1.28	6.34	4.95	0.98	8.07	5.79	
1 784772         AA478542         1.78         8.75         4.92         0.97         8.14           1 47825         R82041         0.06         0.30         4.91         1.91         6.08           2 77545         N56982         0.43         2.11         4.90         1.20         6.72           3 503201         AA14106         0.02         0.08         4.90         5.15         7.27           5 50114         H16743         1.70         8.32         4.90         0.37         7.89           1 504152         AA904969         0.14         0.71         4.89         1.97         7.89           4 7159         AA903384         7.11         34.71         4.89         1.57         7.62           4 8678         AA033284         7.11         34.71         4.88         2.09         5.39           4 8678         AA03328         5.90         28.69         4.81         2.45         5.47           8 80560         N62372         2.16         10.43         4.84         2.45         5.47           8 80560         AA54554         0.01         0.05         4.80         7.91         5.50           3 2547         R43566         AA	828	186132	H39560	0.01	0.05	46.4	8.10	5.74	0.97	
147825         R82041         0.06         0.30         4.91         1.91         6.08           277545         N56982         0.43         2.11         4.90         1.20         8.72           563201         AA114106         0.02         0.08         4.90         1.20         8.72           50114         H16743         1.70         8.32         4.90         0.37         7.29           1504152         AA904369         0.14         0.71         4.89         1.97         8.45           40721         R55750         0.30         1.49         4.88         1.57         7.62           471598         AA035384         7.11         34.71         4.88         2.09         5.39           486787         AA03228         5.90         28.69         4.87         0.13         3.68           280560         N62372         2.16         10.43         4.84         2.45         5.47           809508         AA45554         0.01         0.05         4.83         8.44         5.04           332451         AA064627         2.91         13.99         4.80         7.91         5.50           838568         AAA56931         19.36	960	784772	AA478542	1.78	8.75	4.92	0.97	8.14	5.65	
277545         N56982         0.43         2.11         4.90         1.20         6.72           503201         AA114106         0.02         0.08         4.90         5.15         7.27           50114         H16743         1.70         8.32         4.90         0.37         7.59           1504152         AA904969         0.14         0.71         4.89         1.97         6.45           40721         R55750         0.30         1.49         4.88         1.57         7.62           40721         R55750         0.30         1.49         4.88         1.57         7.62           40721         R403228         7.11         34.71         4.88         2.09         5.39           486787         AA403228         5.90         28.69         4.87         0.13         3.68           290560         NG2372         2.16         10.43         4.81         2.45         5.47           809508         AA454554         0.01         0.05         4.80         0.71         8.39           32517         R43271         0.01         0.05         4.80         7.91         5.68           838568         AAA56931         19.35	961	147825	R82041	90.0	0.30	4.91	1.91	6.08	6.75	
563201         AA114106         0.02         0.08         4.90         5.15         7.27           50114         H16743         1.70         8.32         4.90         0.37         7.89           1504152         AA904969         0.14         0.71         4.89         1.87         6.45           40721         R55750         0.30         1.49         4.86         1.57         7.62           40721         R55750         0.30         1.49         4.88         2.09         5.39           486787         AA043228         5.90         28.69         4.87         0.13         3.68           290560         N62372         2.16         10.43         4.84         2.45         5.47           809508         AA454554         0.01         0.05         4.83         8.44         5.04           382451         AA064627         2.91         13.99         4.80         7.91         6.50           838568         AAA56931         19.35         92.90         4.80         2.12         5.68	962	277545	N56982	0.43	2.11	4.90	1.20	6.72	6.79	
50114         H16743         1.70         8.32         4.90         0.37         7.89           1504152         AA904969         0.14         0.71         4.89         1.97         6.45           40721         R55750         0.30         1.49         4.86         1.57         7.62           471596         AA035384         7.11         34.71         4.88         2.09         5.39           486787         AA043228         5.90         28.69         4.87         0.13         3.68           290560         N62372         2.16         10.43         4.84         2.45         5.47           809508         AA454554         0.01         0.05         4.83         8.44         5.04           322451         AA064627         2.91         13.99         4.80         0.71         8.39           32517         R43271         0.01         0.05         4.80         7.91         5.50           838568         AAA56931         19.35         92.90         4.80         2.12         5.68	983	563201	AA114106	0.02	0.08	4.90	5.15	7.27	2.27	
1504152         AA904969         0.14         0.71         4.89         1.97         6.45           40721         R55750         0.30         1.49         4.86         1.57         7.62           471596         AA035384         7.11         34.71         4.88         2.09         5.39           486787         AA043228         5.90         28.69         4.87         0.13         3.68           290560         N62372         2.16         10.43         4.84         2.45         5.47           809508         AA454554         0.01         0.05         4.83         8.44         5.04           382451         AA064627         2.91         13.99         4.80         0.71         8.39           32517         R43271         0.01         0.05         4.80         7.91         5.50           838568         AAA56931         19.35         92.90         4.80         2.12         5.68	<b>3</b> 6	50114	H16743	1.70	8.32	4.90	0.37	7.99	6.33	
40721         R55750         0.30         1.49         4.88         1.57         7.62           471598         AA035384         7.11         34.71         4.88         2.09         5.39           486787         AA043228         5.90         28.69         4.87         0.13         3.68           290560         N62372         2.16         10.43         4.84         2.45         5.47           809508         AA454554         0.01         0.05         4.83         8.44         5.04           382451         AA064627         2.91         13.99         4.80         0.71         8.39           32517         R43271         0.01         0.05         4.80         7.91         5.50           838568         AAA56931         19.35         92.90         4.80         2.12         5.68           28         AA456931         19.35         92.90         4.80         2.12         5.68	986	1504152	AA904969	0.14	0.71	4.89	1.97	6.45	6.25	
471596         AA035384         7.11         34.71         4.88         2.09         5.39           486787         AA043228         5.90         28.69         4.87         0.13         3.68           290560         NG2372         2.16         10.43         4.84         2.45         5.47           809508         AA454554         0.01         0.05         4.83         8.44         5.04           382451         AA064627         2.91         13.99         4.80         0.71         8.39           32517         R43271         0.01         0.05         4.80         7.91         5.50           838568         AAA56931         19.35         92.90         4.80         2.12         5.68           20         20         20         20         20         2.12         5.68	998	40721	R55750	0.30	1.49	4.88	1.57	7.62	5.46	
486787         AAA043228         5.90         28.69         4.87         0.13         3.68           290560         N62372         2.16         10.43         4.84         2.45         5.47           809508         AAA54554         0.01         0.05         4.83         8.44         5.04           382451         AA064627         2.91         13.99         4.80         0.71         8.39           32517         R43271         0.01         0.05         4.80         7.91         5.50           838568         AAA56931         19.35         92.90         4.80         2.12         5.68           20         20         2.12         5.68         2.12         5.68	298	471598	AA035384	7.11	34.71	4.88	2.09	5.39	7.16	
290560         N62372         2.16         10.43         4.84         2.45         5.47           809508         AA454554         0.01         0.05         4.83         8.44         5.04           382451         AA064627         2.91         13.99         4.80         0.71         8.39           32517         R43271         0.01         0.05         4.80         7.91         5.50           838568         AA456931         19.35         92.90         4.80         2.12         5.68           28         28         28         28         28         28         28	896	486787	AA043228	5.90	28.69	4.87	0.13	3.68	10.79	
809508         AA454554         0.01         0.05         4.83         8.44         5.04           382451         AA064627         2.91         13.99         4.80         0.71         8.39           32517         R43271         0.01         0.05         4.80         7.91         5.50           838568         AA456931         19.35         92.90         4.80         2.12         5.68           28         28         28         28         28         28         28	698	290560	N62372	2.16	10.43	4.84	2.45	5.47	6.59	
382451 AA064627 2.91 13.99 4.80 0.71 8.39 32517 R43271 0.01 0.05 4.80 7.91 5.50 838568 AA456931 19.35 92.90 4.80 2.12 5.68 28	970	803508	AA454554	0.01	0.05	4.83	8.44	5.04 20.04	1.00	
32517 R43271 0.01 0.05 4.80 7.91 5.50 838568 AA456931 19.35 92.90 4.80 2.12 5.68 28	971	382451	AA064627	2.91	13.99	4.80	0.71	8.39	5.32	
838568 AA456931 19.35 92.90 4.80 2.12 5.68	972	32517	R43271	0.01	0.05	4.80	7.91	5.50	1.00	
. 28	973	838568	AA456931	19.35	92.90	4.80	2.12	5.68	9.90	
						28				

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer sell lines.

BNSDGC D < WC = 1153836A2 →

7	7	7	7	7	7	7	2	7	۰ ,		<b>v</b> (	7	7	7	7	7	2	7	7	7	2	۰ ،	4 (	<b>,</b>	7	<b>2</b>	7	7	7	7	7	7	2	7	7	-	
92'9	5.17	5.29	5.41	6.30	5.23	5,54	5.26	5.86	9 6	ţ. !	6.47	5.84 4	5.62	5.79	6.07	5.36	5.76	5.87	5.58	5.36	5.41		D 1	5.65	6.01	5.93	5.55	1.12	5.01	0.56	5.56	5.46	0.55	5.01	5.63	2.63	
5.63	2.84	0.22	5.60	0.31	5.46	5.14	; <b>2</b>	, i	0.00	02.0	5.51	5.60	5.78	5.77	5.61	5.44	5.55	1.07	5.50	5.73	3	6.5	6.03	5.14	5.73	5.12	5.71	5.23	1.78	5.82	5.04	5.12	5.69	5.60	5.56	3660.91	
0.87	5.24	7.74	2.18	6.57	44.	. c	7.50	4.0	7.87 	5.57	0.0	1.50	1.45	1.28	1.08	1.91	1.25	- KG	97.5	) e	) (	9.3g	0.84	1.58	0.57	1.23	26.0	5.88	5.25	5.37	1.13	1.12	5.46	1.01	0.43	0.55	
4.42	4.42	4.42	04.4	70	90: <b>4</b>	) 	0.4.	<b>4</b> .35	4 8	4.33	4.32	4.32	4.28	4.28	4.25	4.24	1 19	4.10		0 4	<del>-</del> -	4.14	4.13	4.12	4.10	4.09	<b>4</b> .08	4.08	4.01	3.92	3.91	3.90	3.90	3.87	3.87	1221.36	C
1.88	0.08	757	1.5	3.5	· 6	<b>\$</b> 1	5.70	0.40	0.42	0.10	1.07	22.74	0.39	0.27	209	11 37		0.02	2 5	11.43	98.0	1.69	9.76	0.57	9.78	0.83	4.55	4.25	7.30	0.07	10.81	2.97	0.0	1.56	7.56	12.21	
0.42	200	<b>7</b> •	- /:	70.61	3 5	ا ا	0.17	0.09	0,10	0.02	0.25	5.27	90	900	3 5	? ?	8 6	OL.7	1.3/	2.75	0.09	0.41	2.36	0.14	2.38	0.20	1.12	9.	1.82	0.02	277	0.76	0.02	0.40	1.95	0.01	
D80765	008377	K063/2	AA398355	190621	AA404338	AA112978	AA884382	AA626247	AA625791	AA443300	AA205320	T70122	A 4.37.746	W451243	100521	9790CN	AA40509Z	N35579	R60949	N50935	AA883755	AA708431	AA487206	R07012	AA633993	AA862814	T54527	T62552	N79230	H79566	AA676598	UB/045	NSR136	H47089	AA045340	N32071	
10/811		657971	726658	110772	758302	530545	1461715	745523	744911	784589	647816	80946	267.783	13/403	4000	508087	814800	272238	42400	281103	1468824	505997	841287	126695	858292	1469211	73659	79726	292388	239661	882483	210517	247587	193420	487141	260168	
000	200	1010	1011	1012	1013																															5 5	

Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least Table 2-10: Prostate

	1.00	0.59	1.33	1.09	1.00	2.22	2.96	2.07	1 1 1 1 1	2.83	1 09	1.04	0.74	0.30	2 47	1.1	1.0	0.25	0.70	0.38	\$. -	1.00	1 16.0	1.10	0.36	2.35	1.26 1	2.37	0.68	0.69	1 1000	0.48	106	-		1 10	2.0
rcell lines.	1675.21	1262.88	1231.52	761.20	730.07	708.49	657.56	586.88	2000.00	213.33 458.05	446.45	440.43	100.10	24.42	241.57	227.03	220.07	218.92	218.05	214.82	212.83	191.25	176.58	175.32	173.57	168.03	162.70	161.00	153.57	149 67	151.46	110 84	4. 6. 64 4. 0. 64	145.80	135.63	133.03	13/.43
les upregua l of 3 cancer	1.63	1.71	0.98	149	2.76	5.5	1 6	C+.2	5.7	c 6	7.07	28.5	5. 5	77.	0.82	2.50	1.79	0.77	0.60	2.48	2.70	25.	1.07	1 75	28.5	20.	1.13	980	8 6		47.7	8°.0	1.48	0.82	5.69	3.00	0.95
ate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line. Celles upreguated of the Cancer Cell lines. 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.	559.28	421 73	411.27	17:1-14	80.407	244.01	23/.61	219.32	197.29	172.23	154.17	149.45	146.97	88.80	80.84	75.00	74.54	73.32	73.13	72.56	72 19	64.60	50 50 50 50	r (c	23.60	36.53	56.40	9 5	94.74	4/.LG	51.03	50.64	50.20	48.61	46.77	46.43	16.12
ic Epithelial ) fold or high	9	6.0	11.7	158.35	2.55	2.45	2.38	2.19	1.97	6.87	6.56	73.56	49.37	69.16	7.42	3.61	8.93	16.59	23.96	1 43	? - c	8.8	S 6	3.72	20.73	18.15	<del>2</del> 5	14.22	3.65	4.65	0.94	7.71	1.05	9.24	32.83	1.25	25 67
mal Prostat 2 of 3 or 10		ro.0	0.02	0.39	0.01	0.01	0.01	0.01	0.01	0.04	9.	0.49	0.34	0.78	0.03	0.05	0.12	0.23	5	3 6	700	2	0.0	0 0 0	0.35	0.31	0.03	0.26	0.07	0.09	0.02	0.15	0.02	0.19	0.70	0.03	1
ines vs. Nor		N47961	N93470	N48913	N30131	R01937	W20486	AA476502	AA486538	AA875888	AA432058	AA456130	AA406070	AA401342	AA461320	AA426212	H11642	A 4 50 8083	500000	N33323	AA399245	H05961	R61395	AA133554	AA002166	AA171784	H16537	AA432278	AA459400	AA481806	AA001219	AA621324	H47076	AA013353	AA398384	AA443140	
cer Cell Li		280291	307328	279613	268115	124567	327495	785337	840937	1325605	784118	796350	743027	742564	706334	769547	47963	2077708	77//80	270603	726439	44007	37986	586845	427693	594683	49131	782353	810959	838877	362278	1048794	193087	360355	726934	796723	
ate Can		<u>\$</u>	1045	<u>8</u>	1047	1048	1049	1050	1051	1052	1053	3 5	1055	1086	25.5	200	9 0	800	200	1061	1062	<del>5</del>	1064	1065	1066	1067	1068	1069	1070	1071	1072	1073	1074	1075	1076	1077	:

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

	0.02 0.03 0.048 0.04 0.04 0.04 0.05 0.03 0.03 0.01 0.01 0.02 0.03 0.03 0.03 0.03 0.03 0.03 0.03	W70264 0.02 N34482 0.03 AA425743 0.48 AA187595 0.02 H08850 0.01 W88779 0.02 W42996 0.10 N64840 0.39 W2064 0.36 W20641788 0.28 AA180453 0.28 AA4877643 0.36 AA4577843 0.06 AA457784 0.00 N23865 0.30 N46845 0.01 AA12867 0.01 N46845 0.00 AA433910 0.07 N66454 0.02 AA433910 0.07
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lated 3 fold or higher in 3 of 3, or 5 fold or higher in at least Table 2-10: Pros

	-	- •	- ·	-	-	-	-	. •	- ,	<b>.</b> .	-	-	-	-	-	-	-	-	_	-	-		· -	-	-	_	_	-	. 🕶	. •	- •	- <b>•</b>	- •		-	-	-	
2.63	1 56	S	7.80	1.17	0.80	1.68	2.46	2 6	2.00	2:30	1.42	1.00	2.23	2.18	0.64	1.17	1.17	77.6	1.45	7.2.0		) c	9.9	78.6	138	1.40		890	1 70	2 6	59.09	0.12	90.1	0.84	2.16	0.46	0.59	
26.20	67:01	73,65	73.28	74.73	74.38	72.44		70.67	71.10	71.45	71.41	70.65	71.20	68.08	69 82	68.34	67.64		7. 50	07.00	09:50	65.41	F8.43	64.73	\$2.50 \$0.50 \$0.50	07.40	62.07	63.23	62.26	59.72	2.06	60.67	58.64	59.31	56.55	57.49	57.17	
00 44	9.6	2.60	1.21	0.75	96.0	3 7	1.20	1.92	1.38	0.41	1.19	2.32	92.0	2.5	, .	0.1	7:07	1.28	1.35	3.00	0.83	<b>1.64</b>	1.16	0.97	0.70	98.0	1.59	81.1	0.58	1.21	1.29	1.10	1.33	0.81	96'0	4	1.24	-
	26.31	25.93	25.76	25.55	25.46	53.10	25.13	25.02	24.82	24.72	79.67	24.66	24.50	24.37	24.2/	23.82	23.68	23.36	22.80	22.78	22.60	22.38	22.28	22.23	22.20	22.17	21.69	21.68	21.17	20.87	20.81	20.63	20.35	20.32	19.89	19 79	19.73	13.0
•	26.72	33.43	92.0	2 6	80.5	2.31	9.24	4.63	449	50 30 30 30		00.01	0.25	0.84	10.63	37.42	10.35	12.38	27.72	6.49	9.38	2.19	14.30	17.18	0.81	8.49	4.05	12.75	59.98	10.68	0.18	1.66	3.00	5.45	2 2	5 5	12.57	<u>=</u>
	1.02	1 20	2 6	500	0.15	0.09	0.37	0.18	;	7 6	2.40	0.65	0.01	0.03	7.0	1.57	0.44	0.53	1.21	0.28	0.42	0.10	0.64	0.77	0.0	0.38	0.19	0.59	1.42	0.51	0.01	0.08	0.15	76.0	7 6	0.43	9. 7.	9
	A A 775041		800+/N	R40057	AA418028	AA425214	44417280	14 KO KO	nene i i	R49462	H97508	N67571	N64774	AA432096	AA884321	R09504	AA011570	AA022668	AA426030	AA417211	AA490264	R92512	AA406220	N66346	W45403	N36389	W74696	AA470073	H89563	NE2079	070700	PAGARA	0.0478		Kokey.	AA885141	N29851	NO4024
	40004	1080/8	59867	27544	767739	773293	711115	21.15.	46303	38280	251961	285682	284569	784154	1466893	127711	429660	364454	757235	731154	8277766	196475	742890	285364	328745	273024	344747	730377	253534	49000	00007	30244	33046	2000	3/901	1466551	259865	440000
	,	1114	1115	1116	1117	1118		8 5	1120	1121	1122	##	1124	1125	1126	1127	1128	00,	5	3 5	5 5	2 3 5 6	2 5	- T	2 2	1437	138	130	5 5	? ;	ויין ל	7611	145	<u> </u>	1145	1146	1147	,

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.

-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-		-	-	-	-	-	-	-	-	<b>~</b> -	-	-	-	-	-	-	-	-	
1.00	1.97	54.90	0.71	0.76	0.38	0.41	1.04	1.41	1.23	1.83	0.50	0.53	1.45	1.64	2.29	0.63	2.70	69.0	1.59	2.06	1.56	0.77	0.74	1.16	0.53	1.30	2.52	44.37	2.06	1.23	1.07	0.38	0.79	26.0	
55.61	53.79	1.00	53.32	54.30	24.12	53.43	51.75	51.86	1.71	49.38	49.70	50.90	0.87	48.99	46.96	49.51	1.50	0.05	46.45	45.77	46.79	47.42	46.41	46.01	46.29	44.76	42.94	1.00	42.61	43.72	42.03	44.40	43.90	42.90	
1.00	1.56	1.10	2.52	0.98	0.38	0.31	1.27	0.55	50.03	1.35	2.13	0.40	49.43	0.80	1.89	0.36	45.76	48.82	1.51	1.72	0.92	1.01	1.38	1.23	1.38	0.93	1.31	1.32	1.79	1.18	2.79	1.07	1.08	1.48	
19.20	19.11	19.00	18.85	18.68	18.29	18.05	18.02	17.94	17.66	17.52	17.44	17.28	17.25	17.14	17.05	16.83	16.65	16.52	16.52	16.51	16.42	16.40	16.18	16.13	16.07	15.67	15.59	15.56	15.49	15.37	15.30	15.28	15.26	15.12	34
0.19	8.40	0.19	0.19	12.10	0.48	14.96	2:32	17.51	7.11	6.75	1.37	3.05	8.13	 48.	6.13	1.60	5.82	3.57	6.03	5.21	3.70	11.22	4.98	14.63	1.63	4.23	5.43	0.16	2.75	8.40	7.61	3.82	8.56	12.92	
0.01	0. <b>1</b>	0.01	0.01	0.65	0.03	0.83	0.13	0.98	0.40	0.39	0.08	0.18	0.47	0.11	0.36	0.10	0.35	0.22	0.37	0.32	0.23	99.0	0.31	0.91	0.10	0.27	0.35	0.01	0.18	0.55	0.50	0.25	9:0	0.85	
AA065042	AA167120	W74725	AA421256	T80832	R43189	AA884397	AA725397	H99075	AA634379	AA620607	AA188710	AA625641	AA418000	AA706935	R60731	R43017	AA884926	N50880	AA412403	N51709	H98676	AA620301	R68409	N40202	AA190825	R51506	AA495938	N75473	A1005321	N66348	W44338	AAB45168	A421170	A083514	
		344618	-	109275	32551	1461725	1342650	262060	743828	1048694	626186	745330	767449	451871	42225	31869	1468260	281003	731422	279102	261989	1030808	137853	276438	627401	38677	768445	239162	1626197	285370	323599	1412504	731091	549157	
1149	1150	1151	1152	1163	15	1155	1156	1157	1158	1159	8	1161	1162	1163	±	1165	1166	1167	±	1169	1170	1171	1172	52	1174	1175	1178	117	1178	1179	1180	1181	1182	1183	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

		6	5.05	15.11	0.85	43.54	96.0	
388373	-	??	3 6	14.83	40	42.32	1.13	
731444	AA412435	0.50	7.34	2,47	1 17	41.82	1.29	
30086	R41782	0.27	6.03	5 4	36	42.02	0.68	
279616	N48302	0.58	8.55	00.41	5 5	40.40	1.92	
280270	N47954	0.53	7.71	14.55		41.08	0.42	
474149	AA911971	0.02	0.34	4.74	2 ;	41.04	0.95	
80242		1.49	21.55	14.50	1.5	50.0	1 16	
00000	A A 241483	0.05	0.78	14.36	0.98	08.04	2 6	
00076			1.87	14.30	1.45	41.37	900	
450375		2 6	75.8	14.22 -	0.76	41.22	0.67	
784242	•	<b>3</b> 5	5	17.99	2.13	1.37	38.45	
25933	R36969	0.01	<del>-</del>	9 6	1.19	39.26	1.48	
42433	R61189	0.15	2.05	0.30		0 11	0.99	
45501	H08397	0.09	1.31	13.90	00.04	5 6	53	
0,48,40		0.58	8.10	13.87	0.91	39.10		
		77.0	10.61	13.85	1.22	39.02	1. 1	
284845		3 6	5	13.84	2.64	2.34	36.52	
744417	_	0.0	ž ;	, c, c,	1 10	39.01	1.27	
328689	W45330	0.70	40.00 40.00	6 F 6 F		39,43	0.85	
342089	W60894	1.08	14.87	13.73	9	39.92	0.39	
38015	R61556	0.17	2.31	13.67	0.69	38.40	2.87	
810511	•	1.09	14.75	13.59	16.1	) to 6	17.0	
2000		0.62	8.43	13.52	0.89	28.80 0	- u	
207907			2.77	13.50	0.86	39.39	67.0	
549035			7.69	13.50	0.89	38.18	1.42	
612782	•	)C'O	5. c	15 27	0.86	38.68	0.57	
730888	3 AA417031	98.0	12.65	13:31	1.42	38.10	0.57	
34901	R45114	0.27	3.60	13.37	<u> </u>	28.03	0.79	
243410	N38992	0.90	12.03	13.35	77	20.00	0.65	
730398	`	<u>5</u> .	13.46	13.30	0.95	5.00	2.48	
		800	0.26	13.29	2.26	41.05	2 9	
7780	•	<b>S</b>	8.68	13.27	1.42	37.10	1.29	
/3005/	•	3 6	40.1	13.26	6.73	38.34	0.72	
283045		65.0	10.7	2 6	1 19	37.93	0.53	
796130	0 AA460967	9. %	4.53	13.22	27.0	38.93	0.11	
591814	4 AA143467	0.09	1.19	13.15	9.45		1,60	
108031		0.22	2.89	13.01	1.76	33.03	33.0	
3		40	14.04	12.99	1.47	36.87	0.00	
/4/000	AA4002/3	3 4	7 09	12.96	1.33	36.05	1.51	
4032073		-	00.					

egulated 3 fold or higher in 3 of 3, or 5 fold or nigher in at least Table 2-10: Prost

9NSCCC D 4 MC - 1153836AC - -

		1	70 67	700	36.14	1.62	-
W84815	0.63	8.15	12.91	78.O	20.00	80.0	
W94591	0.52	99:9	12.84	46. 1	36.41	9 9	· •
T64192	1.07	13.65	12.78	1.20	36.50		- ,-
4A577165	0.33	4.20	12.72	35.38	1.48	05.	- •
AA418988	0.67	8.45	12.60	1.20	35.01	8C. 1	- ,
N75394	0.46	5.81	12.58	<u>.</u> 2	35.31	1.39	<b>.</b> ,
A A053239	0.01	0.13	12.52	1.8	35.57	1.00	-
PK6647	0.21	2.68	12.49	66.0	35.21	1.26	-
NAROON	0.48	90.9	12.49	1.77	33.25	2.43	<b>-</b> -
AAA84317	0.67	8.34	12.47	1.55	35.13	0.71	<b>-</b> .
UEDAGO	0.33	3.96	12.44	35.07	1.49	0.76	-
702102	0.52	6.39	12.37	<u>2</u> .	24.7	0.79	-
2460	200	0 19	12.35	0.66	99.0	35.74	<del>, .</del>
520478	a a	2.21	12.29	1.20	34.01	1.66	-
A068/01		\$ 5	12.20	0.77	34.90	0.95	-
0775M		4 08 4	12.16	1.40	33.59	1.50	-
AAU29428	, t	24.58	12.10	2.01	32.82	1.46	-
77765147	2 4	7.7.	10.01	1,1	33.72	1.28	<del>, .</del>
PA400281	3 4		11.98	0.97	34.33	99.0	-
0018000	5	0 12	11.96	9:1	2.56	32.33	-
060717	0.80	9.57	11.96	0.87	33.75	1.25	-
365384	0.87	10.22	11.73	0.82	33.81	0.58	-
M62780	1,35	15.85	11.70	1.91	32.40	09'0	-
HERORE	80.0	06:0	11.63	1.22	32.62	1.06	<b>-</b>
A A 4 4 0 7 4 5	0.29	3.36	11.63	0.78	32.51	1.61	-
W74471	0.30	3.53	11.62	1.68	31.51	1.68	~
AA862485	0.23	2.61	11.47	33.19	0.28	0.95	-
AA700604	6.42	73.54	11.46	31.50	1.22	1.66	-
T91244	0:30	3.38	11.38	1.20	32.53	0.40	-
AA411682	0.18	2.02	11.35	0.58	33.21	0.27	-
N21233	3	0.41	11.26	1.70	30.58	1.51	-
AA448872	99.0	7.40	11.26	1.20	31.68	06.0	_
H92234	90.0	0.68	11.23	2.84	29.73	1.13	-
AA702840	98.0	4.08	11.23	32.54	0.83	0.32	-
AA469922	0.48	5.33	11.16	1.35	30.72	1.42	-
77	P F O	3	2				

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

•	<b>•</b>		-	-	<b></b>	-	-	-	-	-	-	-	-	-	<b>-</b>	<b>-</b>	-	<b>,-</b> -		-	• •	- •	-	<del>-</del>	-		<b>-</b>	<b>,-</b> -	-	-	-	-	-	-	Ψ-	-		
	1.05	66.0	0.52	0.78	0.63	0.36	0.67	2.58	96.0	0.84	1.80	1.25	1.56	¥.	29.91	60.0	76.0	0.24	1.36	\$	3 6	8	1.60	8	1.00	1.07	0.45	1.21	0.82	1.26	1.36	0.90	1.89	0.88	2.33	06		
	31.28	30.94	31.46	31.48	31.25	34.52	31.48	29.45	101	30.86	20.68	29.36	29.74	29.70	0.06	31.04	20 17	30.56	0000 00000	28.00	29.40	28.01	28.15	28.46	28.58	28.32	29 62	28.06	28.36	27.58	96.96	27 67	26.24	78.75	26.45	20.10	3.7	
	1.1	1.32	1.26	100		57.1	 	5 5	S 65	30.06	6.0	10.0	8 6	1.02	208		0.73	1.65	6/.0	1.01	1.26	1.13	1.20	1.12	1.20	55	3 6	1.32	- v	8. ·	25.	1.09 1.09	1.33	8.	0.49	1.03	1.30	
	11.15	11.09	11.08	20.1.	60.11	11.04	11.00	10.99	10.95	10.89	10.89	10.76	10.75	10.74	10.70	0.00	10.62	10.60	10.52	10.45	10.41	10.40	10.32	40 26	97.01	10.28	10.24	10.20	10.20	10.19	10.01	10.00	9.97	9.92	9.87	9.84	9.84	37
	89 6	2000	60.5	88.11	7.28	3.51	1.56	3.26	2.62	15.93	9.08	2.00	<del></del> ¥.	5.45	2.56	2.63	1.13	4.27	2.53	4.74	6.94	18.36	97. 6	0.50	05.4	0.10	3.44	1.53	4.33	14.09	5.32	3.47	6.03	7.21	4.57	6.60	4.4	
i i	, a	600	0.07	1.08	0.66	0.32	0.14	0.30	0.24	1.46	0.83	0.46	0.12	0.51	0.24	0.25	0.11	0.40	0.24	0.45	0.67	; F	<u> </u>	55.0	4.0	0.01	₹.0	0.15	0.42	1.38	0.53	0.35	0.90	0.73	0.46	0.67	0.45	
		N30328	AA191495	AA188999	AA496949	N33550	R40244	H17038	R42763	T68445	N59278	H99460	R41389	N47214	AA463463	H06525	R51236	AA401482	AA219315	D45402	A A SPORES	AA230301	AA446658	AA412512	AA885140	T50397	AA460289	AA426038	AA453466	N51107	AA609415	AA598943	AA024459	H06380	H96738	W70242	A A 4 59657	
	!	258063		626462	_	_	28774	50582	31478	83358	289725	262313	29251	280413	811785	44387	38517	742616	629968	35308	2000	6/6/89	784177	730294	1466549	72663	795723	773423	795170	282000	743519	898038	364950	44303	251685	344010	705540	200
		1254	1255	1256	1257	1258	1259	1280	1261	1262	1263	1264	1265	1266	1267	1268	1269	1270	27.5	1/7	777	1273	1274	1275	1276	1277	1278	1279	1280	1281	1282	1283	1284	1285	1286	1287	1031	9971

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

-	-	-	-	-	-	-	<del></del>	-	-	-	-	-	-	-	-		-	-	-	-	-	Ψ-	-	-	<b>-</b> -	-	-	-	-	-	-	-	-	-	
0.95	0.81	2.	5.0	1.08	26.83	1.31	0.24	<del>.</del> .	0.43	2.20	<del>1</del> .83	23.15	0.93	0.74	2.61	2.67	0.12	25.05	2.88	1.80	0.49	1.00	0.65	2.23	1.22	1.86	1.78	1.78	1.18	1.23	1.79	99'0	2.72	1.26	
27.38	27.43	26.98	27.37	27.09	0.40	26.66	27.97	26.48	26.99	24.20	26.65	2.81	2.59	26.29	23.30	23.87	27.22	1.00	24.01	25.00	25.75	25.66	25.56	24.09	24.82	24.08	2.38	23.59	24.68	24.09	24.14	25.33	23.64	24.08	
1.16	1.15	1.25	1.1	1.02	1.83	1.01	0.74	1.31	1.25	2.24	0.87	2.58	24.67	1.17	2.22	1.48	0.68	1.90	96.0	1.03	1.55	9,1	1.38	1.09	1.18	1.07	22.82	1.61	2.	1.57	96.0	0.81	0.35	1.26	
9.82	9.80	9.76	9.75	9.73	69.6	99.6	9.65	9.61	9.56	9.55	9.52	9.51	9.40	9.40	9.38	9. 20.	9.34	9.32	9.28	9.28	9.26	9.22	9.20	9.14	9.07	8 <sup>0</sup> .00	8.99	8.99	8.97	8.96	8.96	8.93	8.91	8.87	38
12.27	5.41	5.49	7.41	4.86	1.71	1.53	1.56	6.78	4.97	9.52	4.77	33.20	3.37	6.72	0.72	1.69	0.94	0.09	99.7	1.40	0.19	0.09	3.06	2.38	3.63	3.37	19.69	0.91	6.33	0.48	3.83	8.93	0.87	0.07	
1.25	0.55	0.56	9.76	0.50	0.18	0.16	0.16	. 0.71	0.62	1.00	0.50	3.49	0.36	0.71	0.08	0.18	0.10	0.01	0.82	0.15	0.02	0.0	0.33	0.26	0.40	0.37	2.19	0.10	0.71	0.05	0.43	9:	0.10	0.01	
197599	W79499	AA873089	AA406213	N52876	H11730	H52446	H05626	H89331	AA432141	AA199717	R61871	AA115076	AA459941	N70654	H23555	R39901	H05741	771042	AA127069	AA705858	W69649	T49325	W74633	AA774645	N35603	R41357	AA426113	AA485429	N27272	AA490058	W70074	AA437107	AA156873	AA035144	
121633	347020	1325751	742874	283619	47793	201931	43422	253314	781506	647397	43072	491565	796398	299347	51806	26844	43815	84141	502664	1292207	343871	67397	346642	399319	272288	29424	757265	811050	257608	839903	343977	767352	502527	471859	
1289	1290	1291	1292	1293	1284	1285	1296	1297	1298	1289	1300	1301	1302	1303	900	1306	1306	1307	1308	1309	1310	1311	1312	1313	1314	1315	1316	1317	1318	1319	1320	1321	1322	1323	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithellal Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

-	• •	-	-	-	<del>-</del>	<b>*-</b>	<del>-</del>	-	-	-	-	-	£	-	-	-	<b>~</b> -	<b>-</b>	<b>-</b>	<b>*</b>	<del></del>	<del></del>	<b>~</b>	-	<del></del>	<del>-</del>	<del></del>	<b>-</b>	<b>-</b>	<del>-</del>	<del>-</del>	<del>-</del>	<del>-</del>	<del>-</del>	-
																	0.94																		
																	3 22.84																		
			. •														1.56																		
																	3.53 8.45																		
				•				•						·			0.42																		
	AA459983	H07920	4407028	MEER BO	000000	AA777803	M112003	AA409840 AA608840	AA39030	AA452125	AA778206	AA423318	R157/9	06526N	AA9/4483	A44200F	W46577	4463643	N63453	A A 8 1 2 9 9 6	AA181646	44398262	B42490	A 4047338	AA885471	AA702785	N72150	WE7712	N53534	4448283	AA704255	R45672	R43452	AA416782	
	796448	45578	430814	278900	600017	00000	30300	90400	201069	786550	448379	1/32/10	53377	574407	37283	27,754	204400	811845	286261	1276853	070518	776600	31056	500495	1466418	448014	291129	240898	284220	782847	450574	35503	32381	731337	
	1324	422¥	625	1320	1351	8251	675	<u> </u>	25	1332	1333	138 4	1335	1336	135 155 155	25.	2 5	2 5	<u> </u>	285	2 5	Į ;	2 42	5 5	, 25 25 25 25 25 25 25 25 25 25 25 25 25 2	5 5	136	25.	3 5	300	35.	1355	1356	1357	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines

Table 2-10: Pro

		rostate Cancer Cen Lines.  2 of 3 or 10 fold or higher in at least 1 of 3 cancer cen lines.		5 20	i	2 01 3 01 10 10 10 10 10 10 10 10 10 10 10 10			•	
			•	9	7.31	1.72	18.94	<del>1</del>	<b>-</b> .	
1394 8	823772	AA490240	0.18	2	7.28	1.18	19.21	4	<b></b>	
1395 10	1048283	AA620760	0.43	3.14	27.	1 76	18.29	1.76	<del>-</del>	
	277322	N57475	0.31	2.27	17.7	7	19.38	1.29	-	
	42387	R59990	0.20	1,44	7.26	1,72	19.82	69.0	-	
	307174	N91797	1.14	8.29	7.25	77.1	8	0.54	-	
	417296	W88786	1.03	7.41	7.17	26.0	10.41	1.40	-	
•	399152	AA733105	60.0	0.63	7.15	<b>*</b> 500	10.54	0.81	<b>-</b>	
-	503583	AA131240	0.58	4.11	7.14	7.07	19.61	1.79	-	
	200000	NE4814	0.26	98.	7.12	2.96	16.61	800	-	
•	20400	VATABOZ	0.19	1.32	7.12	20.99	87.0 •	2 45	· •-	
	10000	W/ 400	0.21	1.50	7,11	16.41	2.76	j +	-	
•	415814	W84770		1.74	7.08	0.51	19.69	50.	- 4	
1405	754008	AA478965	C7-0		7.07	1.30	18.84	1.08	_ ,	
1406	39161	R51865	0.22	n i	70	1.41	18.55	1.15	•	
1407	277974	N63436	0.10	0.70	5 6	98.0	19.09	0.95	-	
	729954	AA412046	0.91	6.35	00.7	0.62	18.94	1.36	-	
	813735	AA453795	0.36	2.53	/B.O	18 11	0.81	1.98	-	
	796442	AA459980	2.01	13.98	6.97	16.68	2.00	2.14	-	
	626842	AA191424	44.0	3.06	6.94 6.04	50.5	18.56	1.19	<b>-</b> -	
	281936	N51069	96.0	6.77	5.63	. 4 5. 4	0.20	19.02	<del>-</del>	
	291790	N67861	6.24	43.23	. 6.93 	20.1	17.91	1.36	-	
1414	838818	AA457675	0.35	2.38	6.90	4 +	17.72	1.82	•	
1415	781489	AA432144	0.45	3.11	0.90	300	18.47	o 4	<del></del>	
1416	283703	N50733	0.58	3.88		5 5	18.92	0.76	₩.	
1417	39266	R51872	0.10	0.66	9 6	5 5	18.61	0.99	-	
1418	590120	AA156022	99.0	4.70	20.0	9 9 9	17.58	1.81	-	
1419	27508	R40037	0.24	1.65	0.00	8 6	19.11	0.59	<del>-</del>	
1420	730742	AA435988	0.75	5.74	6.67	6.64	17.15	1.70	-	
1421	213634	H72107	0.85	5.80	6.83	5 5	18.38	0.89	-	
1422	50618	H17516	0.28	1.93	6.83	7 7	18 44	0.93	-	
1423	1030854	AA621750	22.0	4.90	6.81	<u> </u>	18 33	0.91	-	
1424	564847	AA129217	<b>-</b> <b>2</b> 0.	7.07	6.83	9 9	17.77	1.36	<b>-</b> -	
1425	302073	N89783	670	2.00	6.80	97.1	19.47	0.55	-	
1426	730146	AA412477	0.43	2.92	6.80	6.53	17.89	18.1	-	
1427	307660	N92901	99.0	4.43	0.80	5.0 87.0	19.11	0.52	-	
1428	795334	AA454170	0.80	5.43	9.80	2				
					41					

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

1422424 781046 376040 260216 788234 247862 27333 280308 11048586 460398 727275 788240 341081 1055144 796369 341081 361656 335182 816051 1417935 384939 745493 384938

Table 2-10: Prostate

in at least	
s fold or higher	
ate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.	0.86 18.24 2.08 1.53 1.03 1.08 1.28 1.31 1.76 1.04 2.05 1.76 1.30 1.00 2.80 0.72 1.30 1.25 1.25 1.30 1.25 1.30 1.25 1.30 1.25 1.30 1.30 1.25 1.30 1.25 1.30 1.25 1.30 1.25 1.30 1.25 1.30 1.26 1.30 1.25 1.30 1.25 1.30 1.25 1.30 1.25 1.30 1.25 1.30 1.25 1.30 1.30 1.25 1.30 1.25 1.30 1.25 1.30 1.30 1.25 1.30 1.25 1.30 1.25 1.30 1.30 1.25 1.30 1.25 1.30 1.25 1.30 1.25 1.30 1.30 1.25 1.30 1.25 1.30 1.25 1.30 1.25 1.25 1.30 1.25 1.25 1.30 1.25 1.30 1.25 1.25 1.25 1.25 1.30 1.25 1.25 1.25 1.25 1.25 1.25 1.25 1.25
3 fold or highe Il lines.	16.90 0.43 15.24 16.26 17.18 16.71 16.50 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.03 16.
mal Prostatic Epithelial Cell Line: Genes upregulated 3 fold o 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.	1.42 0.49 1.80 1.27 0.85 1.16 0.65 1.16 0.99 1.18 1.14 1.25 1.53 1.01 1.01 1.01 1.01 1.05 1.05 1.05 1.07 1.05 1.05 1.05 1.07 1.05 1.05 1.07 1.05 1.05 1.05 1.07 1.05 1.05 1.05 1.05 1.05 1.05 1.05 1.05
ell Line: Gener rin at least 1 c	6.39 6.39 6.34 6.35 6.29 6.29 6.29 6.29 6.20 6.10 6.11 6.11 6.11 6.13 6.09 6.05 6.05 6.05 6.05 6.05
ic Epithelial C	6.15 7.19 1.88 3.64 6.43 1.245 1.45 1.45 1.45 1.04 7.16 0.70 2.50 0.15 6.12 1.04 7.16 0.16 0.15 6.12 1.04 1.109 0.15 0.15 0.15 0.15 0.15 0.16 0.16 0.17 0.16 0.16 0.17 0.16 0.16 0.17 0.16 0.16 0.17 0.16 0.16 0.17 0.16 0.17 0.16 0.17 0.16 0.17 0.16 0.17 0.16 0.17 0.16 0.17 0.16 0.17 0.16 0.17 0.16 0.17 0.16 0.17 0.16 0.17 0.16 0.17 0.16 0.17 0.16 0.17 0.16 0.17 0.17 0.16 0.17 0.17 0.17 0.16 0.17 0.17 0.17 0.17 0.17 0.17 0.17 0.17 0.17 0.17 0.17 0.17 0.17 0.17 0.17 0.17 0.17 0.17 0.17 0.17 0.17 0.17 0.17 0.17 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.19 0.19 0.19 0.19 0.16 0.16 0.17 0.16 0.17 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.19 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18 0.18
nal Prostati 2 of 3 or 10	1.28 1.12 0.30 0.57 0.85 1.19 0.25 1.15 0.21 0.11 0.02 0.39 0.18 0.47 0.40 0.40 0.40 0.40 0.40 0.40 0.65 0.65 0.65 0.65 0.65 0.65 0.65 0.6
nes vs. Non	AA48270  AA421271  W60288  AA453993  AA465069  R51100  AA406069  RA448855  AA417355  AA16873  AA48855  AA488739  AA495181  AA49181  H10011
ser Cell Li	505454 782838 795254 487383 823578 743025 37404 731198 303023 727137 665033 461420 591540 731050 135900 609530 782888 277631 1669463 277631 1669463 742928 897641 1469857 770989
ate Can	1464 5 1465 1 1467 1 1469 1 1470 1 1471 1 1472 1 1474 1 1479 1 1481 1 14

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

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0.86	0.75	17.23	0.34	1.12	2.48	7.0	<b>T</b> ;	16.74	1.90	0.40	2.85	0.28	13.10	1.22	0.88	0.67	1.27	0.87	0.91	2.69	1.05	14.81	0.56	96.0	0.77	0.18	0.07	1.79	13.64	1.08	1.31	2.88	2.58	14.81	<u>5</u> .	
15.89	15.78	0.01	16.22	15.41	14.53	20:1	15.80	0.54	15.08	0.16	13.14	14.95	3.00	15.23	15.93	15.63	15.07	2.49	15.63	13.52	14.88	99.0	15.61	14.85	16.15	15.29	15.60	14.20	2.35	15.05	14.61	13.08	11.46	1.00	14.64	
1.18	1.32	09'0	1.28	1.27			1.10	0.45	0.72	17.10	1.67	2.42	1.55	1.16	0.79	1.30	1.23	14.09	0.88	1.07	1.36	1.77	1.04	1.37	0.26	1.68	1.44	1.10	1.08	0.88	1.05	1.01	2.86	1.00	1.1	
5.98	5.95	5.95	5.95	5 63	) i	7 7 7	5.91	5.91	5.90	5.89	5.88	5.88	5.88	5.87	5.87	5.87	5.86	5.81	5.81	5.76	5.76	5.75	5.74	5.72	6.72	5.72	5.70	5.70	5.69	5.67	5.66	5.65	5.63	5.61	5.59	
2.05	4.68	3.98	2.17	7.7	7.00 7.00	1.49	4.69	12.14	1.35	2.18	2.11	0.21	37.89	2.08	06.9	6.10	1.46	2.46	2.13	2.71	1.60	4.87	5.93	6.22	24.55	0.32	0.76	3.16	27.34	6.19	2.93	2.33	3.19	90.0	2.90	
¥.0	0.79	0.67		က် (၁)	<b>5</b>	0.25	0.79	2.08	0.23	0.37	0.36	900	4.9	0.35	1.18	5	0.25	0.42	0.37	0.47	0.28	0.85	1.03	1.09	4.29	90.0	0.13	0.55	4.80	1.09	0.52	0.41	0.57	0.0	0.52	,
AA457528	A A 4 D 1 4 D 9	5734CIA	0/9170	AA478928	AA453256	AA775405	AA448825	172235	AA628132	H24329	AA418004	H09616	AA774678	19707N	AA130955	NOSAZE	DAGA7724	H43707	44620027	NAROO1	N49619	R40780	W88999	N23652	AA443958	AA133469	N33366	N50729	AA485653	AA171426	H96605	AA463225	W95118	N33778	W58209	
838500	743184	268446	2007	772938	795339	878097	786107	85840	1055727	51749	767461	46183	856388	298162	588923	247054	R13807	184256	1055636	279583	277785	28106	417461	254564	757190	586796	270681	283702	840404	594806	251517	797048	415191	272951	341095	
1400		3 5	2	1502	1503	1504	1505	1506	1507	809	9 9 9	16.40	5 5		1 1	2 7	# C 4	6 6	2 6 7 7	101	2 4	45.00	1521	1522	1523	1524	1525	1526	1527	152B	15.20	1530	1631	1532	1533	)

Table 2-10: Prostat

ner in at least			-																													•				
tate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least tate Cancer Cell Lines.	+ +		- •	- •	- ,	-	<b>-</b>	-	-	<del></del>	-	-	-	-	-	-	-	-	-	-	-	<del>.</del>	-	-	-	<b>-</b>	<b></b>	-	-	-	-	<del></del>	-	<b>-</b>	<b>-</b>	
lher in 3 of 3, c	1.39	· ·	ŧ ;	\$ 9 5 0	0.43	2.93	0.80	10.87	1.76	25.0	0. <b>5</b>	14.48	0.93	2.71	14.43	79.0	0.98	0.77	1.59	0.69	13.75	10.73	0.14	0.52	\$.	0.22	1.99	0.11	1.81	0.95	1.12	1,40	1.52	12.24	0.60	
ed 3 fold or hig cell lines.	14.07	12.89	0.45	13.78	14.42	12.33	14.57	2.90	1,81	14.66	15.17	1.00	14.52	12.03	1.14	15.22	14.23	14.78	13.47	14.20	1.33	2.97	13.71	15.03	13.09	14.13	11.67	15.56	1.81	14.19	14.02	13.69	12.04	2.43	14.21	
mal Prostatic Epithelial Cell Line: Genes upregulated 3 fold 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.	1.31	1.17	14.87	2.63	1.89	1.47	1.33	2.92	13.10	28	06.0	1			1.02	6.9	0.5g	77.1	0.88	8. 4	1 27	79.0	2.49	0.78	1.86	1.92	2.61	0.59	12.59	20.	9	1.07	2.60	1.48	1.30	
Sell Line: Ger er in at least '	5.59	5.59	5.59	5.58	5.58	5.58	5 57	. v	, u	00.0	0.0	† (i	200	5.52	5.52	5.50	5.48	5,48	5.48	5.47	5.45	0.4. 0.4.	44.9	, 4 4	5.43	6.43	5 A 7	£ 42	5.41	930	);; 30,	5.30	60.5 20.3	5. 3. 0. 3.	5.37	45
tic Epithelial ( ) fold or high	1.02	6.24	1.04	0.37	2.24	17.7	0 60	28.7	57.7	0.42	0.65	5.82	90:0	3.10	0.16	6.39	0.57	3.15	4.04	1.09	4.31	6.45	12.51	0.40	0.22	3.92	0.20	6.33	0.28	6.03	00.4	5.75 0.75	3.29	2.17	3.5	£.
mal Prostal 2 of 3 or 10	0.18	2	: 0	2 2	5 6	3.	4.0	0.53	<u>ਦ</u> ਲ	90.0	0.12	<del>1</del> .6	0.0	0.56	0.03	1.16	0.10	0.57	9.74	0.20	0.79	1.18	2.30	0.04	9. 8	0.72	0.05	8	60	0.0	0.86	0.58	0.61	0.40	2.48	0.27
ines vs. Nori	A A 608075	C 1808044	AA620463	ZODLEN	H06249	AA404356	AA186613	W95414	AA598601	H12338	R28860	N50432	H51549	AA401472	AA401470	AA703147	W73523	AA609284	T96913	H09541	W95409	W92011	H56028	AA447547	AA620472	N90514	AA009763	AA459702	R36989	AA460328	N48078	N47425	AA176607	AA205803	AA496544	H89713
cer Cell Li	80000	•	•	281243			625616	358212	898218	148469	133864	280633	179753	742589	742590	434845	344505	1031552	121409	45922	358200	415264	203544	782611	951101	306121	429773	795585	27098	785788	281756	280672	609062	645332	755904	240273
tate Can	į	1534	1535	1536	1537	1538	1539	1540	154	1542	1543	1544	1545	1546	1547	1548	5 25	45.50	1551	1552	1553	1554	1555	1558	1557	1558	1559	1560	1561	1562	1563	1564	1565	1566	1567	1568

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

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0.39 0.75 0.75 0.15 0.42 1.29 0.82 0.03 0.35 0.39 1.09 0.70 0.66		1.17 13.49 0.99 1.08 1.04 0.65 0.65 0.75 0.75 0.30 12.38 1.37 1.37	13.41 13.90 13.96 14.31 14.30 14.89 14.65 1.66 1.66 1.66 1.84 11.84	1.50 1.13 1.13 0.87 0.61 15.12 0.38 1.43 0.87 0.76 13.96 2.04	
0.25 0.75 0.42 1.29 0.90 1.87 0.35 0.35 0.39 1.09 0.70 0.66		13.49 0.90 1.08 1.04 0.65 0.65 0.75 0.75 0.30 12.38 1.37 1.37	1.23 13.96 14.31 14.89 13.87 14.65 1.66 1.66 11.84 11.84	1.13 0.87 0.61 15.12 0.38 1.43 0.87 0.76 13.96 2.04	
0.75 0.42 1.29 0.90 1.87 3.78 0.03 0.03 0.39 1.09 0.70 0.66		0.99 1.08 1.04 0.65 0.68 0.75 0.75 0.30 12.38 1.37 1.37 2.04	13.90 13.96 14.31 14.89 13.87 14.65 1.66 1.66 11.84 11.84	1.13 0.87 0.61 15.12 0.38 1.43 0.87 0.76 13.96 2.04	
0.15 0.42 1.29 0.90 1.87 3.76 0.35 0.33 1.09 0.70 0.70		1.08 1.04 0.65 0.68 0.62 0.75 0.30 12.38 1.37 1.37 2.04	13.96 14.31 14.89 13.87 14.65 1.66 1.48 11.84 11.84	0.87 0.61 15.12 0.38 1.43 0.87 0.76 13.96 2.04	<del>-</del> -
0.42 1.29 0.90 1.87 3.78 0.35 0.33 1.09 0.70 0.66		1.04 0.65 0.68 0.62 0.75 0.30 12.38 1.37 1.37 2.04 1.40	14.31 0.18 14.89 13.87 14.65 1.66 1.48 11.84 11.84	0.61 15.12 0.38 1.43 0.97 0.76 13.96 2.04	
1.29 0.90 1.87 3.78 0.03 0.03 0.39 1.09 0.70 0.70		0.65 0.68 0.62 0.75 0.30 12.38 1.37 1.37 2.04	0.18 14.89 14.20 1.66 1.66 1.48 13.86 11.84	15.12 0.38 1.43 0.97 0.76 13.96 2.04	-
0.90 1.87 0.82 0.03 0.35 1.09 0.70 0.66		0.68 0.62 0.75 0.30 12.38 1.37 1.40 1.40	14.89 13.87 14.65 1.66 1.48 13.86 14.51 11.84	0.38 1.43 0.97 0.76 13.96 2.04	-
1.87 9.78 0.03 0.03 0.35 1.09 0.70 0.66		0.62 0.75 0.50 0.30 12.38 1.37 0.85 2.04 1.40	13.87 14.20 14.65 1.66 1.48 13.86 14.51 11.84	1.43 0.87 0.76 13.96 2.04	-
3.78 0.82 0.03 0.35 1.09 0.70 0.66		0.75 0.50 0.30 12.38 1.37 0.85 2.04 1.40	14.20 14.65 1.66 1.48 13.86 14.51 11.84	0.87 0.76 13.96 2.04 0.63	-
0.82 0.03 0.35 1.09 0.56 0.70 0.66		0.50 0.30 12.38 1.37 0.85 2.04 1.40	14.65 1.66 1.48 13.86 14.51 11.84	0.76 13.96 2.04 0.63	<b>-</b>
0.03 0.35 1.09 0.56 0.70 0.66 1.00	_	0.30 12.38 1.37 0.85 2.04 1.40	1.66 1.48 13.86 14.51 11.84	13.96 2.04 0.63	-
0.35 0.39 1.09 0.56 0.70 0.66	_	12.38 1.37 0.85 2.04 1.40	13.86 13.86 11.84 11.84	2.04 0.63	-
2.14 0.39 1.09 0.56 0.70 0.66	_	1.37 0.85 2.04 1.40	13.86 15.41 13.64	0.63	-
0.39 0.56 0.70 0.66 0.100		2.04 1.40	16.41 48.11 18.12		-
0.56 0.70 0.70 0.66		1.40	4.5. 4.6.	0.49	-
0.56 0.70 0.66 1.00		1.40	13.61	98.	-
0.70 0.66 1.00		14.20	2	0.83	-
1.00		2	1.16	0.47	-
1.00	47 5.28	14.23	0.22	1.38	-
		1.12	13.86	0.83	-
		1.10	13.60	<del>.</del>	-
0.56		1.26	13.15	<b>1.3</b>	-
0.02		0.49	0.41	14.83	┯.
1.79		0.59	13.78	<del>1</del> .34	-
1.08		0.33	12.98	2.39	-
	23 5.23	1.88	12.77	<del>7</del> .	-
4.0		1.87	12.97	0.84	-
0.22		0.13	15.02	0.51	
0.18		0.41	12.55	2.66	-
0.48	5.20	1.26	11.43	2.91	-
0.11		0.63	14.87	60.0	-
0.52		1.54	12.74	1.28	-
0.70		14.21	0.93	0.40	-
2.59	39 5.18	1.16	12.97	1.40	-
1.11		0.73	12.99	1.80	-

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

1,1792/2   AM672279   2.49   12.87   5.17   0.92   1,1798   AA608609   2.49   12.87   5.17   0.92   1,1798   AA608609   2.49   12.87   5.16   1.80   1,170278   AA608602   0.69   3.54   5.16   1.80   1.80   1,170278   AA608602   0.51   2.63   5.16   1.80   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.12   1.1			7.0	1.38	5.17	1.17	12.54	<del>.</del> 8.	-
(1031726         AAA008544         2.74         3.90         5.16         1.37         12.96           (1031726         AAA085034         0.76         3.54         5.16         1.80         13.19           (1031726         AAA086029         0.51         2.63         5.16         1.80         13.19           (1030727         AAA17815         0.59         3.04         5.15         1.17         13.22           341065         W58191         0.52         2.68         5.15         1.67         12.49           341065         W58191         0.52         2.68         5.14         1.92         2.33           233172         H7035         0.42         2.14         5.13         10.92         2.33           105184         AA62865         0.63         3.24         5.13         10.92         2.45           106189         AA         1.11         5.71         5.13         10.73         2.45           106189         AA         1.23         5.13         10.73         2.45         1.15           27927         AA40860         1.11         5.71         5.13         10.73         1.15           27927         AA60860         1.03 <td>14/38/2</td> <td></td> <td>4 6</td> <td>12.87</td> <td>5.17</td> <td>0.92</td> <td>13.61</td> <td>96.0</td> <td>•</td>	14/38/2		4 6	12.87	5.17	0.92	13.61	96.0	•
193179   AA00554   U.76   S.79   S.16   S.16   S.16   S.16   S.15   S.16   S.15   S.16   S.15   S.	742898	•	6 · · · ·	5 6	, Y	1.37	12.96	1.16	_
63081         R16259         0.09         5.24         5.10         4.42           6470278         AA417915         0.51         2.63         5.16         0.43         14.23           76744         AA417915         0.52         2.68         5.15         1.12         13.22           341065         AA417915         0.52         2.68         5.15         1.67         12.49           341065         AA428659         0.63         3.22         5.14         1.92         2.33           781447         AA428659         0.63         3.22         5.14         1.32         13.03           78147         AA428659         0.63         3.22         5.14         1.32         13.33           1030764         AA608974         0.71         3.64         5.13         10.73         2.45           136189         R92281         1.88         10.18         5.13         10.73         2.45           136187         AA408860         0.43         2.19         5.13         10.73         1.76           27927         AA408860         0.17         0.87         5.13         10.73         1.74           27927         AA40860         0.13         2.	1031736		e (	ה ה ה ה	, r	1.80	13.19	0.50	_
1470278         AAB66029         0.51         2.63         5.10         0.22           76744         AAA17915         0.59         3.04         5.15         1.12         1.32           341065         W68191         0.52         2.68         5.15         1.17         12.49           331065         W68191         0.52         2.68         5.13         1.09         1.33           331066         W68191         0.52         2.68         5.13         1.09         1.33           105196         R62281         0.71         3.64         5.13         1.09         1.33           105196         R62281         1.11         5.71         5.13         1.08         1.33           105197         AA20086         0.27         1.36         5.13         1.07         1.75           279278         AA01777         2.39         12.23         5.13         1.07         1.75           279278         AA010777         2.39         1.223         5.13         1.07         1.75           279278         AA010777         2.39         1.23         1.13         1.75           279278         AA460864         1.09         5.13         1.07	53081		69.0	\$ C	5 A	0.43	14.23	0.82	_
7674.4 AA417915 0.59 3.04 3.13 1.11 341065 W88191 0.52 2.68 5.15 1.15 233712 H79035 0.42 2.14 5.14 10.92 2.33 1030764 AA608974 0.71 3.64 5.13 0.80 13.30 1030764 AA608974 0.71 3.64 5.13 0.80 13.30 1292121 AA70765 0.77 1.36 5.13 10.73 1.75 1292121 AA70765 0.77 1.36 5.13 10.73 1.75 1290649 AA407777 2.39 12.23 5.12 11.35 2.00 279278 N88590 0.43 2.19 5.11 1.25 13.31 279278 AA408084 1.09 5.54 5.10 0.66 13.45 840786 AA408084 1.09 5.54 5.10 1.21 1.25 279278 AA408084 0.43 2.18 5.07 1.51 13.26 279278 AA408083 0.45 2.26 5.06 0.60 1.34 812233 AA45096 0.45 2.26 5.01 1.34 11.90 279279 AA408083 0.02 0.43 1.14 5.02 2.55 11.25 257746 N30577 0.82 2.63 5.01 1.70 1.21 257747 N2089 0.85 2.76 5.00 1.21 1.19 257746 N30597 0.82 2.63 5.01 1.70 1.24 257746 AA408083 0.01 0.05 5.01 1.30 257747 AA40390 0.45 2.26 5.01 1.30 257748 N30597 0.82 2.63 5.01 1.70 1.24 257749 AA432075 1.61 7.99 4.97 1.344 0.13 257759 AA432076 0.74 7.99 4.99 0.73 13.08 257650 N67510 0.74 5.69 4.99 0.73 13.08 257651 AA432076 0.74 7.99 4.99 0.73 13.08 257652 AA445004 0.76 2.23 4.99 0.73 13.09 257634 AA432076 1.14 5.69 4.99 0.73 13.09 276539 AA445004 0.74 1.17 4.94 0.43 13.57 276539 AA445044 0.75 1.71 4.94 0.43 13.57			0.51	2.63	0. 10 14 15	+ +2	13.32	29.1	_
341065 WiSHIST 0.52 2.68 5.15 10.01 233712 H79035 0.42 2.14 5.14 10.92 2.33 781447 AA428659 0.63 3.22 5.14 1.32 13.03 781447 AA428659 0.63 3.22 5.14 1.32 13.03 195189 R92281 1.98 10.18 5.13 10.73 2.45 196189 R92281 1.98 10.18 5.13 10.73 2.45 1252121 AA0707615 0.27 1.36 5.13 10.73 1.75 490649 AA070771 0.29 12.23 5.12 11.35 2.00 279278 NASSOG 0.47 0.87 2.19 5.11 1.25 13.11 279276 AA40389 0.83 4.24 5.10 0.66 13.43 278976 NASSOG 0.45 2.18 5.07 1.51 13.26 278977 AA40389 0.83 4.24 5.10 1.21 13.05 278978 AA45096 0.45 2.25 5.06 0.60 13.49 278572 NASSOG 0.45 2.25 5.04 1.69 11.45 257746 N30597 0.82 4.11 5.02 2.60 10.93 278572 NBG178 0.65 2.76 5.00 1.31 11.65 257746 N30597 0.82 2.63 5.01 1.70 12.41 257746 N30597 0.85 2.76 5.00 1.31 1.70 257746 N30597 0.45 2.25 5.04 1.69 1.34 257746 N30597 0.45 2.25 2.63 5.01 1.70 12.41 257746 N30597 0.45 2.25 2.63 5.01 1.30 257746 N30597 0.45 2.26 2.00 1.31 1.20 257746 N30597 0.45 2.26 2.00 1.31 1.30 257746 N30597 0.45 2.26 2.00 1.31 1.30 257746 N30597 0.41 2.40 0.45 1.34 0.43 257747 0.4004946 0.76 2.77 1.34 0.43 25774 0.4004946 0.76 2.27 1.44 0.43 0.43 25774 0.4004946 0.76 2.40 0.98 2.40 25774 0.4004946 0.76 2.40 0.98 2.40 2576 2.40 0.98 2.40 0.73 1.35 25774 0.4004946 0.76 2.40 0.98 2.40 25774 0.4004946 0.76 2.40 0.98 2.40 25774 0.4004946 0.76 2.40 0.40 0.40 25774 0.4004946 0.76 2.40 0.40 0.40 25774 0.4004946 0.77 0.41 25774 0.4004946 0.77 0.41 25775 0.4004946 0.77 0.41 25779 0.4004946 0.77 0.41 25779 0.4004946 0.77 0.41 25779 0.4004946 0.77 0.41 25779 0.4004946 0.77 0.41 25779 0.4004946 0.77 0.41 25779 0.4004949 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.404 0.4		AA417915	0.59	9.0 <del>8</del>	<u>0</u>	9 : ·	12.40	128	_
233712         H70035         0.42         2.14         5.14         10.92         2.33           78147         AA428659         0.63         3.22         5.14         1.32         13.03           78147         AA608974         0.71         3.64         5.13         0.80         -13.91           196189         R92261         1.88         10.18         5.13         10.73         2.45           196180         R92221         1.8         10.18         5.13         10.73         2.45           196182         AA60864         1.11         5.71         5.13         10.73         2.45           1292712         AA707615         0.27         1.36         5.13         10.73         1.75           279276         AA408604         1.09         5.43         2.19         5.11         1.25         1.75           279276         AA408604         1.09         5.43         5.10         1.51         1.36           279276         AA408604         1.09         5.43         5.10         1.51         1.36           289356         N45260         0.43         2.18         5.10         1.51         1.37           278012         N45177 <td>1610 341065</td> <td>W58191</td> <td>0.52</td> <td>2.68</td> <td>5.15</td> <td>1.0/</td> <td>5 to 0</td> <td>9 40</td> <td>-</td>	1610 341065	W58191	0.52	2.68	5.15	1.0/	5 to 0	9 40	-
78147 AA42869 0.63 3.22 5.14 1.32 13.03 103076 AA608974 0.71 3.64 5.13 0.80 13.21 103076 AA608974 0.71 3.64 5.13 10.73 2.45 196189 R92281 1.88 10.18 5.71 5.13 10.73 2.45 1292121 AA707615 0.27 1.36 5.13 10.73 1.75 490649 AA101777 2.39 1.2.23 5.12 11.35 2.00 279278 N48590 0.43 2.19 5.11 1.25 13.11 279278 N48590 0.43 2.19 5.11 1.25 13.11 279278 N48590 0.43 2.19 5.10 1.21 1.35 288936 N487089 0.83 4.24 5.10 1.51 1.35 278806 N487109 0.43 2.18 5.06 1.14 1.32 278812 AA458096 0.45 2.25 5.06 1.14 1.32 287857 AA458096 0.45 2.25 5.06 1.14 1.32 287857 AA458096 0.45 2.25 5.06 1.14 1.32 287857 AA458096 0.45 2.25 5.01 1.14 1.32 287858 N62893 0.52 2.63 5.01 1.70 1.21 285778 N68178 0.82 4.11 5.02 2.55 11.42 287858 N6789 0.82 4.11 5.02 2.55 11.42 287859 N67579 1.14 5.69 5.01 1.30 282785 AA018893 0.01 0.05 5.01 1.70 1.21 285680 N67570 1.14 5.69 4.99 0.73 13.08 285580 N67570 1.14 5.69 4.99 0.73 13.08 28529 AA045804 0.24 1.17 4.94 0.43 28529 AA045804 0.24 1.17 4.94 0.96 2.40		H79035	0.42	2.14	5.14	10.92	2.33	E1:7	• •
1030754         AA608974         0.71         3.64         5.13         0.80         13.91           1030754         AA608974         0.71         3.64         5.13         0.073         2.45           196189         R92Z81         1.88         10.18         5.13         10.73         2.45           276827         AA707615         0.27         1.36         5.13         10.73         1.75           490649         AA101777         2.39         1.223         5.12         11.35         2.00           279278         N48590         0.43         2.19         5.11         1.25         13.11           279278         NA62806         0.17         0.87         5.10         0.66         13.43           289350         NA67806         0.17         0.87         5.10         0.66         13.43           289350         NA67809         0.83         4.24         5.10         0.66         13.43           278912         NA460983         0.35         1.78         5.06         1.21         13.06           278912         NA460983         0.35         1.78         5.06         0.60         13.49           27812         NA460983         <		A4428659	0.63	3.22	5.14	1.32	13.03	1.07	
196169			17.0	39.5	5.13	08.0	- 13.91	0.68	_
1901 89         NAZZEI         110         51.3         1.08         132.0           279827         NAGO76 1.11         5.71         5.13         1.073         1.75           1292121         AAAV07615         0.27         1.36         5.13         10.73         1.75           490649         AA407777         2.39         12.23         5.12         11.35         2.00           279278         NA8590         0.43         2.19         5.11         1.25         13.11           280326         NG2696         0.17         0.87         5.10         1.51         13.26           280326         NG2696         0.17         0.87         5.10         1.51         13.26           742607         AA400389         0.83         4.24         5.10         1.51         13.26           728036         NA41039         0.43         2.18         5.07         1.24         13.28           796123         AA400893         0.83         4.24         5.06         1.14         13.28           796123         AA400893         0.85         4.31         5.06         0.50         1.47           287746         NA55096         0.45         1.29 <t< td=""><td></td><td></td><td></td><td>10 18</td><td>5.13</td><td>10.73</td><td>2.45</td><td>2.21</td><td>_</td></t<>				10 18	5.13	10.73	2.45	2.21	_
27/92/17         NAD270         1.11         1.25         5.13         1.75           1292/12/1 AA707615         0.27         1.36         5.13         10.73         1.75           490649         AA401777         2.39         12.23         5.12         11.35         2.00           490649         AA401777         2.39         12.23         5.10         0.66         13.43           840786         AA400389         0.83         4.24         5.10         0.66         13.43           289356         N49109         0.43         2.18         5.06         1.21         13.26           279806         N49109         0.43         2.18         5.06         1.24         12.47           279806         N49109         0.43         2.18         5.06         1.24         13.48           279806         N49109         0.43         2.18         5.06         1.14         13.28           279806         N49109         0.43         2.18         5.06         1.14         13.28           279806         N491075         0.45         2.25         5.06         1.14         13.28           283122         N45509         0.45         2.25         <			} <del>-</del>	5.71	5,13	1.08	13.20	1.10	_
1292121         AA707615         0.27         1.30         5.12         11.35         2.00           490649         AA101777         2.39         12.23         5.12         11.35         2.00           490649         AA101777         2.39         12.23         5.12         11.35         2.00           279276         AA436084         1.09         5.54         5.10         0.66         13.43           289356         N62696         0.17         0.87         5.10         1.51         13.26           279806         NA5296         0.43         2.18         5.07         1.55         12.47           279806         NA4109         0.43         2.18         5.07         1.55         12.47           279806         NA4109         0.43         2.18         5.07         1.25         12.47           279807         NA45086         0.43         2.18         5.07         1.25         12.47           279807         NA45086         0.43         2.18         5.06         1.14         13.28           28122         AA456086         0.45         2.25         5.06         0.50         1.14           257746         N30597         0.82			=		. t	10 73	1.75	2.91	_
490649         AA101777         2.39         12.23         5.14         1.25         13.11           279276         NA8590         0.43         2.19         5.11         1.25         13.11           240786         AA486084         1.09         5.54         5.10         0.68         13.43           240786         AA486084         1.09         5.54         5.10         1.51         13.26           72567         AA400389         0.83         4.24         5.10         1.21         13.26           72560         AA400389         0.83         4.24         5.10         1.21         13.26           279806         N49109         0.43         2.18         5.07         1.55         12.47           279806         N49109         0.43         2.18         5.06         1.14         13.28           301842         N91175         2.45         1.23         5.06         1.14         13.28           301842         N91750         0.45         2.25         5.06         0.60         13.49           283122         AA456096         0.45         2.25         5.06         2.60         1.69           27776         N26933         0.52 <td>1616 1292121</td> <td></td> <td>0.27</td> <td>1.30</td> <td>2 5</td> <td>11 35</td> <td>200</td> <td>2.00</td> <td>•</td>	1616 1292121		0.27	1.30	2 5	11 35	200	2.00	•
279276         NA8590         0.43         2.19         5.11         1.25         15.11           840786         AA486084         1.09         5.54         5.10         0.66         13.43           288936         NGZ696         0.17         0.87         5.10         1.51         13.26           288936         NGZ696         0.17         0.87         5.10         1.51         13.26           742607         AA400389         0.83         4.24         5.10         1.21         13.06           729806         N43109         0.43         2.18         5.07         1.55         12.47           279806         N43109         0.43         2.18         5.06         1.14         13.28           796123         AA460963         0.35         1.78         5.06         2.38         11.90           796122         AA45096         0.45         2.25         5.06         2.38         11.90           278572         N66178         2.84         14.29         5.04         1.69         11.42           257746         N30597         0.82         2.41         5.02         2.55         11.42           257779         N26993         0.52	1617 490649		2.39	12.23	21.c	S: -		86.0	
840786         AA486084         1.09         5.54         5.10         0.66         13.43           288936         NG2696         0.17         0.87         5.10         1.51         13.26           288936         NG2696         0.17         0.87         5.10         1.51         13.26           742607         AA400389         0.83         4.24         5.10         1.21         13.06           278806         N49109         0.43         2.18         5.06         1.14         13.28           301842         N91175         2.45         12.39         5.06         1.14         13.28           301842         N91775         2.45         1.78         5.06         2.38         11.90           283122         N45223         0.85         4.31         5.06         0.60         13.49           281293         AA456096         0.45         2.25         5.05         2.60         10.93           27774         N30597         0.82         4.11         5.02         2.55         11.42           25774         N30597         0.82         2.63         5.01         1.70         12.41           257779         N5693         0.50			0.43	2.19	5.11	1.25	13.1	3 6	•
288936         NG2696         0.17         0.87         5.10         1.51         13.26           742607         AA400389         0.83         4.24         5.10         1.21         13.06           742607         AA400389         0.83         4.24         5.10         1.21         13.06           279806         N49109         0.43         2.18         5.07         1.55         12.47           301842         N91175         2.45         12.39         5.06         1.14         13.28           301842         N94523         0.35         1.78         5.06         2.38         11.30           283122         N45223         0.85         4.31         5.06         2.60         10.93           812293         AA456096         0.45         2.25         5.05         2.60         10.93           27779         N2693         0.82         4.11         5.02         2.55         11.42           257779         N26993         0.52         2.63         5.01         1.70         12.41           257779         N26993         0.05         2.76         5.01         1.21         11.91           25660         N67819         0.52			1.09	5.54	5.10	0.66	13.43	C7:1	- •
742607         AA400389         0.83         4.24         5.10         1.21         13.06           279806         N49109         0.43         2.18         5.07         1.55         12.47           279806         N49109         0.43         2.18         5.06         1.14         13.28           301842         N91175         2.45         12.39         5.06         1.14         13.28           301842         N91175         2.45         1.78         5.06         1.14         13.28           301842         N95123         0.85         4.31         5.06         0.60         13.49           283122         N455096         0.45         2.25         5.05         2.60         10.93           278572         N66178         2.84         14.29         5.04         1.69         11.65           257746         N30597         0.82         4.11         5.02         2.55         11.42           257779         N26993         0.52         2.63         5.01         13.03         1.00           290680         N6769         0.05         5.01         1.24         0.13           290680         N6756         0.20         0.38			0.17	0.87	5.10	1.51	13.26	0.52	_ ,
279806         N49109         0.43         2.18         5.07         1.55         12.47           301842         N91175         2.45         12.39         5.06         1.14         13.28           301842         N91175         2.45         12.39         5.06         1.14         13.28           796123         AA460963         0.35         1.78         5.06         2.38         11.90           283122         NA5223         0.85         4.31         5.06         0.60         13.49           281293         AA45696         0.45         2.25         5.05         2.60         10.93           27734         NX0597         0.82         4.11         5.02         2.55         11.42           257779         NX0593         0.52         2.63         5.01         1.70         12.41           257779         NX0593         0.52         2.63         5.01         1.70         12.41           25660         NG7619         0.55         2.76         5.01         1.21         11.91           2567779         NX0593         0.52         2.63         5.01         1.21         11.91           25660         NG7619         0.56			0.83	4.24	5.10	1.21	13.06	1.01	
201842         N91175         2.45         12.39         5.06         1.14         13.28           796123         AA460963         0.35         1.78         5.06         2.38         11.90           796123         AA460963         0.35         1.78         5.06         2.38         11.90           283122         NA5223         0.85         4.31         5.06         0.60         13.49           278572         N66178         2.84         14.29         5.04         1.69         11.65           257746         N30597         0.82         4.11         5.02         2.55         11.42           257746         N30597         0.82         4.11         5.02         2.55         11.42           257776         N26993         0.52         2.63         5.01         1.70         12.41           250680         N67619         0.55         2.76         5.01         1.21         11.91           290680         N67619         0.55         2.76         5.01         13.03         1.00           285680         N67670         1.14         5.69         4.99         0.73         13.08           285680         N67570         1.61			0.43	2.18	5.07	1.55	12.47	121	
251225         AA460965         0.35         1.78         5.06         2.38         11.90           283122         NA4523         0.85         4.31         5.06         0.60         13.49           283122         NA455096         0.45         2.25         5.05         2.60         10.93           812293         AA455096         0.45         2.25         5.04         1.69         11.65           27746         N30597         0.82         4.11         5.02         2.55         11.42           257779         N26993         0.52         2.63         5.01         1.70         12.41           257779         N26993         0.52         2.63         5.01         1.70         12.41           250680         N67619         0.55         2.76         5.01         1.21         11.91           250680         N67619         0.55         2.76         5.00         1.21         11.91           119289         T98056         0.20         0.98         5.00         1.31         12.76           119289         T98056         0.20         0.98         5.00         1.31         13.08           285680         N67570         1.14			2.45	12.39	5.06	1.14	13.28	0.75	
283122         NA4503605         3.506         0.60         13.49           283122         NA455096         0.45         2.25         5.06         2.60         10.93           812293         AA455096         0.45         2.25         5.04         1.69         11.65           278572         N66178         2.84         14.29         5.04         1.69         11.65           257746         N30597         0.82         4.11         5.02         2.55         11.42           257779         N26993         0.52         2.63         5.01         1.70         12.41           250680         N67619         0.55         2.76         5.00         1.21         11.91           250680         N67619         0.55         2.76         5.00         1.21         11.91           119283         T98056         0.20         0.98         5.00         1.31         12.76           285680         N67570         1.14         5.69         4.97         13.44         0.13           784130         AA432075         1.61         7.99         4.97         1.24         12.45           345118         W72290         0.76         2.23         4.96 </td <td></td> <td></td> <td>, c</td> <td>1.78</td> <td>5.06</td> <td>2.38</td> <td>11.90</td> <td>0.88</td> <td>•</td>			, c	1.78	5.06	2.38	11.90	0.88	•
255122         N49223         CCS         2.25         5.05         2.60         10.93           812203         AA455096         0.45         2.25         5.04         1.69         11.65           278572         N66178         2.84         14.29         5.04         1.69         11.65           257746         N30597         0.82         4.11         5.02         2.55         11.42           257779         N26993         0.52         2.63         5.01         1.70         12.41           250680         N67619         0.55         2.76         5.00         1.21         11.91           119289         T98056         0.20         0.98         5.00         1.31         12.76           285680         N67570         1.14         5.69         4.99         0.73         13.08           285680         N67570         1.61         7.99         4.97         13.44         0.13           784130         AA432075         1.61         7.99         4.97         13.44         0.13           42829         AA465904         0.76         2.23         4.96         0.78         12.45           42829         AA466904         0.24			3 6	4.33	5.06	0.60	13.49	1.08	•
812293         AA4550040         1.59         11.65           277346         N30597         0.82         4.11         5.02         2.55         11.42           257746         N30597         0.82         4.11         5.02         2.55         11.42           257779         N26993         0.52         2.63         5.01         1.70         12.41           257779         N26993         0.62         2.63         5.01         1.70         12.41           290680         N67619         0.55         2.76         5.00         1.21         11.91           119289         T98056         0.20         0.98         5.00         1.31         12.76           285680         N67570         1.14         5.69         4.99         0.73         13.08           285680         N67570         1.61         7.99         4.97         13.44         0.13           784130         AA432075         1.61         7.99         4.97         13.44         0.13           428238         AA004946         0.76         2.23         4.96         0.78         12.45           42829         AA456904         0.24         1.17         4.94         0.43			3 4	2.25	5.05	2.60	10.93	1.62	•
2778572         N66178         2.04         17.23         2.55         11.42           257746         N30597         0.82         4.11         5.02         2.55         11.42           257779         N26993         0.52         2.63         5.01         1.70         12.41           257779         N26993         0.52         2.63         5.01         13.03         1.00           290680         N67619         0.55         2.76         5.00         1.21         11.91           119289         T98056         0.20         0.98         5.00         1.31         12.76           280580         N67570         1.14         5.69         4.99         0.73         13.08           285680         N67570         1.61         7.99         4.97         13.44         0.13           784130         AA432075         1.61         7.99         4.97         13.44         0.13           428238         AA004946         0.76         2.23         4.96         0.78         12.45           428238         AA456904         0.24         1.17         4.94         0.43         13.67           42824         AA456904         0.24         1.17 <td></td> <td>_</td> <td>2</td> <td>14.30</td> <td>25</td> <td>1.69</td> <td>11.65</td> <td>1.78</td> <td>•-</td>		_	2	14.30	25	1.69	11.65	1.78	•-
257746         N30597         U.O.Z.         2.63         5.01         1.70         12.41           257779         N26993         0.52         2.63         5.01         1.70         12.41           280785         AA018683         0.01         0.05         5.01         1.303         1.00           290680         N67619         0.55         2.76         5.00         1.21         11.91           119283         T98056         0.20         0.98         5.00         1.31         12.76           285680         N67570         1.14         5.69         4.99         0.73         13.08           784130         AA432075         1.61         7.99         4.97         13.44         0.13           784130         AA432075         1.61         7.99         4.97         13.49         0.13           782829         AAA004946         0.76         3.75         4.95         1.05         12.45           813242         AA456904         0.24         1.17         4.94         0.43         13.67           81324         AA456904         0.24         1.17         4.94         0.43         13.67			<b>t</b> 6		5.02	2.55	11.42	1.09	•
257779         N26993         0.52         2.03         3.04         1.00           362795         AA018683         0.01         0.05         5.01         13.03         1.00           290680         N67619         0.55         2.76         5.00         1.21         11.91           119283         T98056         0.20         0.98         5.00         1.31         12.76           285680         N67570         1.14         5.69         4.99         0.73         13.08           784130         AA432075         1.61         7.99         4.97         13.44         0.13           784130         AA432075         1.61         7.99         4.97         13.44         0.13           784130         AA432076         0.45         2.23         4.96         0.78         12.45           345118         W72920         0.45         2.23         4.96         0.78         12.80           428298         AA004946         0.76         3.75         4.94         0.43         13.67           813242         AA456904         0.24         1.17         4.94         0.43         13.67           400         0.96         0.96         0.240			70.0	- <b>(</b>	20.3	1.70	12.41	0.93	•
362795 AA018683 U.01 U.02 U.03 U.03 U.03 U.03 U.03 U.03 U.03 U.03			70.0	20.2	, r	13.03	1.00	1.00	•-
290680 N67619 0.55 2.70 3.00 1.31 12.76 119289 T98056 0.20 0.98 5.00 1.31 12.76 119289 T98056 0.20 0.98 5.00 1.31 12.76 13.08 285680 N67570 1.14 5.69 4.99 0.73 13.08 13.08 245116 W72920 0.45 2.23 4.96 0.78 12.45 12.45 1428298 AA004946 0.76 3.75 4.95 1.05 12.80 13.67 13.74 0.43 13.67 13.67 13.74 0.98 2.40			5	0.00		7.7	11.91	1.89	
119289 T98056 0.20 0.98 5.00 1.51 13.08 285680 N67570 1.14 5.69 4.99 0.73 13.08 13.08 245130 AA432075 1.61 7.99 4.97 13.44 0.13 12.45 345116 W72920 0.45 2.23 4.96 0.78 12.45 12.45 1.25 4.95 1.05 12.80 13.24 0.74 1.17 4.94 0.43 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 13.67 1			8	2.70	8 6		12.76	0.93	•
285680 N67570 1.14 5.69 4.59 0.73 0.73 0.73 0.73 0.73 0.73 0.73 0.73			8	9.0	00.5	<u>.</u> 6	80 4	1.15	•
784130 AA432075 1.61 7.99 4.97 13.44 0.13 345118 W72920 0.45 2.23 4.96 0.78 12.45 428298 AA004946 0.76 3.75 4.95 1.05 12.80 813242 AA456904 0.24 1.17 4.94 0.43 13.67			1.14	5.69	4.99	5.5	9.5		•
345118 W72920 0.45 2.23 4.96 0.78 12.45 '428298 AA004946 0.76 3.75 4.95 1.05 12.80 813242 AA456904 0.24 1.17 4.94 0.43 13.67	•		1.61	7.99	4.97	13.44	0.13	1.33	
428298 AA004946 0.76 3.75 4.95 1.05 12.80 813242 AA455904 0.24 1.17 4.94 0.43 13.67			0.45	2.23	4.96	0.78	12.45	<u>7</u>	
813242 AA456904 0.24 1.17 4.94 0.43 13.67	•		0.76	3.75	4.95	1.05	12.80	0.98	•
2.40		•	0.24	1.17	4.94	0.43	13.67	0.73	•
7577R H77047 DG C 1707R DS 2707R		•	920	27.18	4.94	96.0	2.40	11.45	•

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

BNSDCC/D + WC \_\_0153836A2\_\_\_5

589861 346987 743058 811899								•
346997 743058 811899	AA158117	0.68	3.33	4.94	0.78	12.51	<u>.</u>	-
743058 743058 811899	M70446	2.28	11.24	4.93	0.21	14.40	0.20	-
1642 /43056 1643 811899		2 6	2.47	4.93	1.19	12.16	4.	-
	•	3. 6	11 69	4.93	1.24	11.25	2.29	-
000110	-	5 -	6.40	4.92	1.30	12.51	96:0	-
	•	3 4	0.81	4.92	1.09	10.94	2.74	-
	•	2 5	5.51	4.92	2.39	10.12	2.25	-
	•	4 6	. vo c	4 91	0.82	13.48	0.44	-
		0.18 0.48	3.35	16.91	1.14	12.92	0.66	-
_	•	9 6	1 + 5	16.9	1.98	12.52	0.22	-
		5.0	1::	4 90	66.0	13.24	0.48	-
_	•	<b>5</b>	S .	06 4	14.47	0.18	0.05	_
•	AA101632	0.24	0 4	8 6	109	1.08	12.48	_
1652 49203		0.24	1.13	D 6	55.	12 12	96.0	_
1653 280763	N50563	<b>9</b> .0	0.19	88.	1.36	7	7	-
1654 731393	AA421054	0.27	1.31	4.88	1.60	06: I	77.6	•
Ī		1.1	5.56	4.88	1.59	12.60	S :	- ,
		1.40	6.82	4 87	2.38	0.46	11.76	_ `
_		0.37	1.82	4.86	1.34	11.70	1.55	_
		. <del>2</del>	8.97	4.86	0.79	12.99	0.81	-
		2.68	13.01	4.86	1.09	11.94	1.55	-
		0.13	0.65	4.86	1.07	11.25	2.25	-
		0.0	0.05	4.86	1.00	1.00	12.57	-
26296		0.43	2.06	4.85	1.10	12.19	1.25	-
•	T96032	2	3.08	4.84	1.16	12.34	<del>7</del> .	-
1/7011 500		1,7	8.56	4.84	1.76	1.07	11.69	-
•			80.5	4.84	1.0	11.69	1.78	-
	NO4003	5 6	35.0 35.0	4.83	2.	12.71	0.13	_
	•	3 5	20.5	4 83	0.83	1.93	11.72	-
		7 6	9 6	4 87	1.36	11.63	1.48	-
_		0.0	5.55	1 C	1.06	11.76	0.74	-
	_	2.68	76.21	. 4 10. 8	28.0	11.57	1.92	-
670 824358	AA489681	0.48	2.35	4.02		37.04	7.4	•
743416	AA609334	1.12	5.41	4.82	0.99	12.76		- •
672 289666	97777N	9.0 8	0.21	4.82	1.07	11.91	1.48	- '
•	•	3.28	15.82	4.82	10.78	1.98	.68 83.	-

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

	<b>-</b> -	-	-	•		-	-	-	- 1	-	-	-	-	. •	- ,	_	<b></b>	<b>-</b>	-	-	. +			<b>-</b>	<del>-</del>	<del>-</del>	-	-		-	-	-	-	-	-	-	-	-	-		
	1.32	1.36	11 14		3	10.93	0.38		67.0	1.79	0.71	1.56		0.00	0.60	1.12	1.55	0.57	1.27	08.0	9. (	0.42	0.88	1.16	1.07	0.60	12.09	0.62	1.17	0.41	11.48	1.84	1.18	1.08	10.93	0.94	12.11	141	1 00	<u>.</u>	
	12.21	1.38	3 3	1.32	0.33	1.75		12.91	13.94	12.05	12.67	5.5	10.55	12.47	13.16	11.33	11.93	12.85	44.28	07.11	12.29	11.88	12.35	11.65	0.73	11.61	1.53	12.86	11.17	12.67	0.35	11.08	11.59	11.34	1.15	11.22		5 5	10.1	05.11	
	0.92		11.55	1.89	2.18	1 67	70.	1.06	0.10	0.47		<b>\$</b>	2.16	0.91	0.47	1.17		200	<b>8</b> (	1.57	1.23	1.82	0.87	1.26	12.24	1.83	0.40	25.0	9	880	2 12	060	÷ 5	S 65	2	1.12	3 5	1.67	1.32	1.45	
	č	. <del>4</del>	4.79	4.79	92.	4.70	4.78	4.78	11.1		4.7.	4.77	4.75	4.75	4.74	,	4.74	4.73	4.72	4.71	4.71	4.73		07:4	n 60 +	90.4	00.4	00.4	70.4	70.4	3 4	6.4	10.4	16.4	00.4	4.60	9.4	4.59	4.58	4.58	49
2 of 3 or 10 told to 10.		3.96	0.04	26.60	20.00	6.99	260.67	50	5.5	21.07	0.65	5.60	0.34		04.0	3.45	4.38	6.86	1.32	2.04	683	5 6	0.47	3.89	3.20	2.71	0.92	5.40	5.18	3.00	5.43	9.34	0.82	5.04 4	2.73	4.90	9.37	103.69	1.74	1.38	
2 of 3 or 10		0.82	5	5 1	5.35	1.46	54.50		0.42	4.41	41.0	1 17		9	0.72	0.73	0.92	1.45	0.28	670	?	1.20	0 9 9	0.83	0.68	0.58	0.20	1.15	1.1	0.64	1.17	2.01	0.18	1.09	0.59	1.06	2.04	22.57	0.38	0.0	;
		AAA64419		R35665	H27379	4598567		070C0M	AA005358	AA278759	A A O 57520	2001000	AASAAA	AA609422	AA459857	AA779048	N49098	N59295	NEBOOR	07600N	AA634482	AA452578	AA454654	AA461086	AA431975	9600£N	W20462	N91811	AA195651	H05777	AA011597	AA417895	T89084	N39590	AA401436	AA877815	AA486445	A & B 73050	A 6 4 1 2 4 B 5	60471400	W73022
		700400		137017	163174	202105	CE1060	295483	428202					743536	795810	858761	77977	280704	10107	C9/67	743901	788519	\$11907	796170	782208	256907	327480	306513	665148	43936	429681	752770	22541	277063	743193	1161564	842927	4475055	14/0003	2000	345318
		į	1674	1675	1676		16//	1678	1679	1680	3	1681	1682	1683	1684	1685	200	9 6	1991	1688	1689	1690	1691	1692	1693	1694	1695	1608	1897	1698	1699	1700	107	1702	4704	404	100	601	1706	1707	1708

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least אינים א

-	-		-	-	-	- •		<b>.</b>	<b>-</b>	-	<del></del>	<b>-</b>	_	-	-	-	-	-	· •	- •	- •	<b>.</b> .	<b>.</b>	_	<b>-</b> .	-	-	-	-	-	<del>-</del>	_	-	-	_	<b>-</b>	
1.70	1.05	1.26	1.25	0.28	0.46	7	<b>3</b> .	11.60	2.11	2.26	2.26	0.51	0.44	0.98	<b>1</b> 9.1	1.50	1.29	0.76	2.75 11.46	Q# (1	0.39	1.39	0.71	1.00	2.45	0.87	0.63	2.19	<del>2</del> .	0.77	1.31	1.03	0.21	1.12	0.90	0.48	
11.01	11.06	11.38	11.52	12.54	5 9	12.10	1.8	06:0	0.97	10.76	10.02	12.14	12.07	11.61	10.60	11.03	10.50	0.00 82.00 83.00	0	00.	12.03	10.20	12.07	0.61	10.04	11.55	11.78	10.29	10.92	11.30	10.60	11.12	12.55	11.25	11.45	0.39	
0.97	53.	86.0	28.0		6.73	1.05	1.23	<del>2</del> .	10.45	0.52	1.26	0.88	1.01	0.90	1.24	. E6 C	9.0	9 7	112	1.00	<b>1</b> .04	1.80	0.58	11.75	0.86	0.91	0.91	0.82	<u>2</u> .	1.21	<del>2</del> .	1.10	0.49	0.88	0.89	12.36	
4.56	4.55	75 7	7 7	ţ ;	<b>₹</b>	4.53	4.53	4.51	4.51	4.51	4.51	4.51	4.51	4.50	4 49	9 7	7	64.4	4.49	4.49	4.48	4.46	4.45	4.45	4.45	4.4	4.44	4.43	4.43	4.43	4.42	4.42	4.42	14.4	4.41	4 41	ř
3.82	2.28	2 7	4. c	9.79	<b>4</b> .00	4.48	4.06	4.46	1.66	2.99	1.40	3.80	204	0.50	50.0	6.63	71.7	3.23	4.55	0.04	6.20	8.04	2.59	2.85	2.32	3.08	1.74	9.64	1.90	1.88	1.11	4.58	4.12	1.25	5.27	3,5	0.40
28	5 6	) i	0.63	0.82	0.88	0.99	0.60	0.99	0.37	990	0.34	84	0.00 4.50	? ;	2 4	Ç. (	0.48 8	0.72	1.01	0.01	1.38	1.80	0.58	20.	0.52	0.69	0.39	0.15	0.43	0.42	0.25	1 1	86.0	0.28	5	? ;	76.1
E 830014	Secuen	AA808430	H97514	AA454844	W92798	AA702788	AA487046	AA703117	H17022	00406200	A A R34470	A 45,005	AA404000	AA38624/	K45/00	W79525	AA666087	AA101833	N22210	N63768	W86832	AA464861	AA682549	AA777098	R98407	NS1574	H11631	T58430	4444781	N59870	A A 085759	A A 4 1 2 4 0 0	A 4 4 7 7 9 4 7	B30014	10000	17077	ZCALEN
305743	300243	143531	251212	809976	418400	448024	841153	434828	50781	77877	773803	20001	100540	(20037)	2040	347066	859382	485549	254468	293013	416889	789383	431255	377898	200656	280431	48060	75644	705740	284803	488776	730447	773812	24895	430742	2001	4/2807
,	1709	1710	171	1712	1713	1714	1715	1746	2 (	2 2	9 5	n (	02/1	1271	1722	1723	1724	1725	1726	1727	1728	1779	27.0	1731	1732	1	1734	1735	477	1737	12.4	2 5	927	27.4	;	76/1	1743

Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least Table 2-10: Prost

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.

-	<del>-</del>	-		-	-		-	-	-	-	<b>-</b>	-	-	-	-	<del>-</del> -	-	-	-	-	-	-	<b>-</b>	_	-	-		_		-	-	_	_	_	
0.52	1.12	1.07	0.49	0.81	12.16	0.91	0.89	0.88	26.0	0.80	10.41	1.75	1.05	1.35	0.82	0.76	0.00	0.31	10.18	0.53	10.22	0.23	1.26	11.28	0.28	9.76	0.91	1.03	1.25	0.72	95.0	0.55	0.91	0.62	
10.99	0.35	10.39	0.51	0.02	0.15	10.52	10.31	11.10	10.02	10.33	1.00	10.14	10.78	10.06	10.57	10.58	10.32	10.47	1.18	10.08	0.02	11.66	10.38	0.27	11.02	10.39	10.21	10.27	10.00	10.15	10.61	10.42	10.16	10.43	
1.08	11.10	1.10	11.57	11.74	0.25	1.12	1.35	0.54	14.	1.25	0.93	0.41	0.47	0.89	0.89	0.93	0.95	1,43	0.85	1.60	1.93	0.28	0.52	0.50	0.82	96.0	96.0	0.75	0.79	1.16	0.93	0.96	0.80	0.80	
4.20	4.19	4.19	4.19	4.19	4.19	4.18	4.18	4.17	4.13	4.13	4.11	4.10	4.10	4.10	4.09	4.09	4 09	4.07	4.07	4.07	4.06	4.06	4.06	4.04	4.04	40.4	4.02	4.02	4 01	4 01	4.00	3.98	3.96	3.95	25
3.49	5.55	1.75	6.49	2.67	38.53	3.65	4.43	0.35	3.11	4.41	0.04	1.03	2.80	2.10	2.74	2.71	1.43	0.13	2.12	6.21	8.26	3.16	14.17	3.89	3.07	6.18	1.77	1.08	0.90	2.00	8.94	4.1	2.21	6.19	
0.83	1.33	0.45	1.55	0.64	9.21	0.87	1.08	90.0	0.75	1.07	0.01	0.25	0.68	0.51	0.67	99.0	0.35	9.03	0.52	1.53	2.04	0.78	3.49	0.98	0.76	1.53	14.0	0.27	0.22	0.50	2.24	0.29	0.56	1.57	
AA453465	N91003	AA412446	N47941	W95082	AA620995	AA421001	W81117	AA449686	AA455130	AA470079	T50995	AA778533	W80791	AA455358	AA707753	R05660	AA425749	AA479912	H20046	AA447561	AA487146	AA292054	AA187287	AA701933	R15891	AA443823	AA399633	H08560	AA621478	AA136071	N78903	AA194189	R37780	AA159613	
795168	306066	730100	280217	415145	1056172	731298	415413	785760	809863	730406	76647	1048872	347268	812222	412983	29397	773392	772916	172785	782617	841253	725365	625465	435573	53122	784064	729924	45542	1055278	490071	300024	665738	27072	593218	
1779	1780	1781	1782	1783	1784	1785	1786	1787	1788	1789	1790	1791	1792	1793	1794	1795	1796	1797	1798	1799	1800	1801	1802	1803	1804	1805	1806	1807	1808	1809	1810	1811	1812	1813	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

1.00 0.70 0.22 0.68 10.05 0.14 0.74 0.74 0.74 0.02 0.02
10.07 0.79 11.14 10.25 0.95 10.29 10.20 10.26 10.26 10.08 0.12 0.12 0.12
0.76 10.25 0.46 0.87 0.74 1.30 0.88 0.68 0.78 0.78 0.79 0.79 0.70 0.70 0.70 0.25 0.56 10.29
3.60 3.60 3.60 3.60 3.60 3.70 3.70 3.60
2.48 0.05 0.98 3.36 9.98 0.77 10.36 0.92 0.93 0.42 4.06 24.47 1.73 0.11
0.63 0.01 0.25 0.85 2.55 0.20 0.24 0.19 0.05 0.05 0.05 0.04 0.04
AA150401 AA479906 AA479056 AA479313 H08598 AA4783169 R60717 AA426087 R60705 AA426087 R60705 AA476867 AA625890 AA176867 AA148735
491690 772904 784105 1031279 76433 45783 1455671 42115 795736 449438 757238 757238 41903 1466416 744944 609155
1814 1815 1816 1817 1820 1821 1822 1823 1825 1825 1825 1825 1825

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostastic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines

Order	IMAGE Clone ID	Gen Bank Accession Number	MPM1(PrEC)	Ave Cancer Cell Lines Values	Fold(AveCell line/MPM1)	Fold(MPN2/MP M1)	Fold(MPM2/MP Fold(MPM272/M # of Cancer M1) PM1) Cell Lines	Fold(MPM272/M PM1)	# of Cancar Cell Lines
	757165	AA443950	0.01	5.71	591.67	7.33	66.37	1701.31	က
. ~	592707	AA160606	0.01	1.66	234.30	8.10	680.66	14.13	ന
, m	220851	H95633	0.01	1.28	196.56	8.10	570.75	10.82	က
•	782804	AA448186	0.01	1.81	180.70	8.88	526.08	7.15	ო
· vo	1343732	AA725564	0.01	1.12	112.48	13.45	301.13	22.85	ო
, ε	784296	AA447079	0.01	99.0	65.30	45.84	106.60	43.45	က
. ~	796569	AA460463	100	0.35	63.5/	16.10	138.91	35.70	ო
. 00	1558655	AA976561	90'0	4.36	54.32	97.06	17.92	47.97	m
6	743016	AA406036	0.01	0.50	50.47	15.70	123.80	11.93	ന
. <del>6</del>	39442	R51617	0.10	5.02	49.93	99.52	14.7	5.49	ო
F	33076	R44048	0.80	36.21	45.35	56.36	29.19	50.51	ന
12	626390	AA189106	0.07	3.16	45.16	91.62	20.97	22.90	8
5	128083	R09747	0.01	4.0	44.34	25.03	101.07	6.93	ო
<u> </u>	812170	AA456035	0.01	0.30	37.18	16.25	80.79	14.51	es
5	50288	H17888	0.01	0.32	36.79	12.25	87.45	10.68	m
16	243181	H94482	0.01	0.36	35.62	5.91	88.28	12.68	m
17	256911	N30098	0.01	0.25	34.44	20.19	73.72	9.41	က
8	1030613	AA608832	0.01	0.33	33.18	5.81	82.72	11.01	60
6	487861	AA045436	0.02	0.59	32.57	29.44	39.44	28.82	က
8	270626	N33331	0.01	0.32	31.52	8.48	9.21	76.87	က
21	291057	N72115	0.23	6.85	29.19	24.48	38.30	24.80	6
22	1455835	AA863292	0.01	0.40	29.18	9.10	30.24	48.14	9
23	108378	1777729	0.01	0.41	29.13	7.73	36.47	43.19	n
24	1391682	AA789328	0.02	0.44	28.83	66.5	65.78	14.72	က
25	263014	H99813	0.63	17.34	27.31	19.07	21.59	41.29	e
56	303180	N92764	0.01	0.27	27.21	13.82	36.04	31.77	n
27	42415	R60981	0.12	3.11	26.71	9.46	65.35	5.30	က
					•				

Table 2-11; Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

768056	AA418903	0.01	0.16	26.24	7.36	35.03	36.33	e .
20001	LINGRAB	900	1.21	25.96	33.86	30.16	13.87	r>
05004	108010	0.48	12.53	25.93	38.89	13.26	25.64	m
131 161	H02294	9	200	26.84	81.3	58.23	14.11	60
38344	R49555	0.0	0.24	7 70	12.08	52.20	12.48	က
41595	R59556	.0.0	<u>.</u> 6	65.55	24 ± 55	13.61	26.60	ю
430092	AA009840	0.40	00.01	24.00	46.38	16.26	10.88	n
1049033	AA778675	0.38	00.00 00.00	16.42	92 52	10.09	32.50	က
884790	AA629838	5 6	0.75 4.00	73 07	44.37	17.55	9.88	n
23345	R39191	6.9	0.0	23.01	8 11	34.50	26.43	က
284714	N63057	D:0	2 G	22.74	38.20	16.96	11.55	က
365149	AA025142		70.0	00 00	5.07	7.89	49.72	က
418094	W90036	5 6	17.0	20.02	18.18	18.45	24.62	က
51581	H22824	5	0.70	20.42	5.36	33.42	22.42	က
270975	N32542	0.0	0.50	Pr. 00	20 00	30.57	9.30	က
51916	H22563	0.12	2.40	20.21	70.2	11.138	40.27	ю
40010	R54036	0.01	0.19	19.30	, t	24.35	20.16	ю
1467161	AA883187	0.02	0.31	16.24	10.25	00 80	14 13	es
488431	AA047441	0.17	2.95	17.83	10.45	06.02	27.0	er.
435099	AA701315	0.01	0.21	17.36	7.52	33.02		۰ «
67440	T49355	 8.	33.78	17.25	25.95	8.91	10.00	3 6
280155	N47012	9.0	0.63	17.15	5.64	11.78	¥ 5	ი •
454083	AA676998	0.01	0.17	16.89	8.16	10.67	31.84	o (
3.475.RB	W81432	0.0	0.17	16.56	7.84	8.92	32.94	က
289606	10101 N26769	0.01	0.21	16.54	6.72	36.98	5.91	ന
0000346	14802063	50	0.16	16.35	10.47	5.09	33.48	m
330330	W92203	5 6	60°0	16.34	9.38	30.25	9.40	9
25.55	11.0400		11.31	16.17	5.81	15.89	26.82	က
25/5/55	W\$37 (7	2 6	0.33	15.94	21.48	14.07	12.25	က
80070	0/2501		000	15.63	6.61	33.79	6.49	က
015/0/	AA416343	20.00	8 6	15.00	33.97	5.66	5.36	က
011111	0.520	2.0	0.35	14.92	5.14	24.61	15.00	m
200007	767676	3 6	3.0	14.61	£.1	16.46	22.27	က
08707	0.0437442	5 5	0.14	14.54	5.64	32.86	5.13	n
25915	R39951	0.0	0.14	14.20	9.65	6.51	26.44	က
1031446	AA609189	9.0	0:0	14.14	5.44	31.74	5.25	က
				2				
				,				

PCT/US01/02318

Table 2-11; Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

32109	R42780	0.09	1.25	14.14	31.21	6.86 29.82	5.35 5.12	<b>6</b> 1 61
0820	R42600	50.0	15.63	13.71	10.6	9.76	20.74	ы
U35/30	AA626607	7.7	16.52	13.65	13.47	11.85	15.62	<b>6</b> 7
20000	A 4 2 8 6 1 7	10.0	0.17	13.58	7.33	15.99	17.41	m .
20,000	AA016225	0.0	0.13	13.43	9.24	8.92	22.12	ი (
5089	W93379	0.65	8. B.	13.24	8.23	12.04	19.45	n (
68409	AA885478	0.14	1.91	13.23	22.16	9.42	8.12	<b>~</b> (
1103	H18217	0.02	0.23	13.14	14.72	17.85	6.86	m c
44050	N34042	0.01	0.17	12.88	5.65	21.72	11.26	n •
97807	AA598531	0.29	3.56	12.37	21.95	7.81	1.37	, e
\$41	AA844550	0.01	0.12	12.30	5.26	7.58	24.06	<b>~</b> ~
86172	AA262211	1.18	14.39	12.18	5.65	13.80	17.09	o (
2070FC	A A B 0 9 0 0 0	900	0.31	11.73	10.15	8.33	16.72	ю ·
1080	H67712	8	0,45	11.30	5.08	7.88	20.94	<b>с</b> э (
0000	0.0463433	0	60.0	10.70	11.07	8.10	12.92	ო .
60700	HEROTA	13	13.51	10.64	11.48	13.18	7.26	က
430028	A678205	0.74	2.	10.34	20.21	5.77	5.03	ო -
76457	AA041107	600	0.32	9.78	5.69	8.56	15.10	<b>~</b>
23667	AA704652	8 0	0.16	9.57	11.04	10.28	7.37	ო
3220	M47113	1 2	10.61	9.52	11.98	10.79	5.79	က
	N00770	4	1.40	9.48	5.53	11.42	11.50	m
66000	1000E44	2 2	2.04	9.46	6.47	14.40	7.50	က
8000	AA/0264	1 5	10.53	9.44	5.48	13.22	9.62	ဗ
09300	M24042 A A O 4 5 6 5 5	0.37	3.43	9.32	9:36	11.24	7.35	က
13733	H04789	0.11	1.03	9.24	10.06	12.44	5.24	es (
34617	H77727	0.07	0.60	9.16	12.99	9.66	5.84	m
77884	H46176	0.16	1.4	9.15	5.38	8.45	13.62	m
07056	AA598817	0.15	1.40	9.13	13.21	8.47	5.71	<b>6</b> 0
24004	W/40672	100	0.11	8.92	7.38	12.47	6.90	ന
78247	750699	0.01	0.05	8.82	9.14	7.21	10.11	ო
524527	44187351	0.56	4.90	8.72	5.30	9.17	11.70	က
101184	AA137072	149	12.90	8.67	6.69	13.78	5.53	m
26273	AA520999	0.45	3.90	8.62	5.02	10.99	9.84	က
42906	R60019	0.19	1.57	8.31	9.78	8.33	6.83	m
				ო				

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

n n	6	· (	· •	m	ო	n	n	, m	က	ы	ന	ო	ო	n	က	ဗ	ю	2	7	7	7	7	7	2	8	2	7	2	2	7	7	7	64	7		
8.51	900	6.03	7.97	1.50	7.97	9.46	8.78	7.98	6.19	5.60	6.63	5.79	5.08	5.21	5.52	5.77	5.40	7.96	7.05	1917.92	5.67	6.10	9.57	9.33	7.73	1.35	5.84	146.14	3.45	105.59	12.64	13.76	7.41	99.88		
11.10	20.0	6.22	7.09	5.10	6.65	6.44	7.63	7.83	5.36	5.02	5.78	5.67	5.73	6.93	7.41	563	5.01	2966 06	2506.90	61 14	1831.96	1516.35	1171.33	944.19	841.07	674.66	302.09	140.40	249.63	150.46	239.88	229.57	203.18	95.72		
5.27	11.86	8.56	7.77	40.9	7.30	10.8	, <del>,</del> ,	; u	3 5 5	9.45	, t		14.7		S 6	7 G	n (	n 0	2. 4	 	7.7	 	1 6	2. 2.	; ; ;	9. £	60 6	9 0	, r	5 6	, c	5.00	, c	5 6	Š.	
8.30	7.87	7.61	7.61	1 :01	5 6	15.7	05.7	61.7	7.05 7.05	C (8)	6.69	6.30	6.13	6.10	6.05	5.99	5.63	5.20	991.80	838.64	660.46	613.07	508.28	393.88	319.30	283.26	228.80	103.67	95.78	86.03	85.65	85.17	81.38	71.26	65.49	4
7.83	0.54	68.9	9 6	5.53	cr.0	3.90	0.12	18.03	3.05	0.72	2.07	47.65	7.33	16.66	29.59	0.07	1.94	0.12	9.92	8.39	9.14	9.00	5.08	3.94	3.19	2.83	1.70	<del>2</del>	1.93	2.66	6.83	0.51	0.82	0.71	5.16	
46.0	20.0	, i	٥ ز.	0.73	0.05	0.53	0.02	2.51	0.43	0.10	0.31	7.49	1.20	2.73	4.89	0.01	0.3	0.02	0.01	0.01	0.01	0.01	0.01	0.01	0.01	0.01	0.01	0.01	0.05	0.03	0.08	0.01	0.01	0.01	90.0	
AA425755		AA455938	R96941	AA608568	N30372	H29227	AA477400	AA663310	R06944	AA701645	R28287	H02231	R10284	AA425302	AA459318	R93875	H08785	R52085	N74995	AA009615	R60995	N57483	AA433885	N64379	AA099431	AA429904	AA46446	AA424900	H56453	AA071503	AA485214	AA459917	W80611	AA876021	AA419486	
772383	2000	813271	200402	950690	260035	52704	740620	853368	126650	433576	134719	151055	128947	773073	814306	275871	45882	40299	299559	365551	42627	277327	773640	290213	510631	781157	781036	768226	203878	366039	815794	795627	347182	1161830	746152	
ć	9	66	5	5	102	103	\$	505	8	107	80	8	110	Ξ	112	1 1	4.	. <del>.</del>	1 1 1	117	. 18	119	52	121	122	123	124	125	126	127	128	129	130	131	132	:

Table 2-11; Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

7	2	7	7	2	7	7	7	7	7	7	7	7	2	7	8	7	2	7	23	2	7	7	2	7	7	3	7	7	7	2	3	7	7	2	
11.76	0.73	4.30	70.45	123.70	0.72	1.93	5.39	6.47	12.17	91.51	7.45	6.23	3.16	11.05	19.33	5.11	2.18	51.22	6.23	5.61	7.60	55.73	81.28	5.06	0.70	11.03	11,15	71.43	52.45	1.00	63.66	10.31	9.84	6.05	
168.10	173.62	170.43	4.69	51.92	88.50	154.39	150.40	1.37	139.36	56.58	128.30	120.65	121.33	109.53	99.20	106.34	105.49	52.03	89.38	0.12	81.20	34.54	7.97	76.38	<b>3</b> 9.6	70.25	69.04	1.36	27.83	72.00	11.75	66.04	62.71	68.60	
3.73	6.88	6.29	104.24	1.03	78.84	5.36	45.0	144.61	0.63	0.91	2.24	3.54	5.62	4.59	4.81	3.08	5.72	0.93	1.79	87.92	2.89	0.77	1.00	1.61	72.37	0.74	1.21	8.33	0.69	5.77	2.28	0.59	4.23	2.05	
61.20	60.41	60.34	68.79	58.88	56.02	53.90	52.11	50.82	50.72	49.67	45.99	43.48	43.37	41.72	41.11	38.18	37.79	34.73	32.46	31.22	30.56	30.35	30.08	27.65	27.63	27.34	27.13	27.04	26.99	26.26	25.90	25.65	25.59	25.57	5
0.61	0.83	1.60	26.93	2.87	12.41	0.28	2.21	6.09	1.10	13.02	09.0	0.81	0.43	0.42	0.41	0.38	0.38	8.56	1,12	2.62	0.70	7.89	0.30	5.88	16.16	0.27	0.27	0.20	14.38	0.26	14.15	0.39	0.29	1.90	
0.01	0.0	0.03	0.45	9.09	0.22	0.0	900	1.20	0.02	0.26	0.01	0.02	0.0	0.0	0.01	0.0	0.0	0.25	0.03	90.0	0.02	0.26	0.01	0.21	0.58	0.01	0.01	0.01	0.53	0.01	0.55	0.02	0.01	0.07	
W94289	AA757754	AA461522	R23270	AA251784	W73144	AA086475	R40231	44437224	R44741	AA504844	AA49120	AA855158	R37395	AA437128	AA810225	AA062985	AAB57748	R91566	R59473	H29290	AA401434	AA485052	AA099369	R49439	R90934	R15740	AA460274	AA443624	H50655	AA425158	N47445	N52767	AA452139	R40018	
346942	395708	795858	131452	684879	344589	562811	30207	757435	33827	825740	785849	1389018	28498	757368	1367900	366209	1435339	196544	37823	49858	743189	815861	489656	38347	167205	53039	796549	771294	194318	769000	280699	283379	787856	27605	
133	2	135	85	137	138	65	9	1		143	. <u>1</u>	14.5	146	147	148	149	150	151	152	153	<u>3</u>	55	156	157	158	159	160	161	162	163	2	165	991	167	

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

77 (	4 (	N (	N (	۷ (	7 (	2	2	~	2	7	2	7	2	7 (	2	7	~	7	~ (	74 (	7	7	7	7	<b>~</b>	~	7	7	7	7	7	7	7	73	
16.63	5.46	8.05	11.62	31.35	45.50	11.25	62.76	43.13	3.46	48.25	25.34	31.76	11.14	53.51	54.20	23.77	3.08	47.62	17.48	28.27	20.17	50.17	45.41	25.26	49.74	46.16	1.08	43.06	5.54	10.71	6.22	5.54	16.22	8.77	
54.64	67.88	64.40	60.05	41.17	26.82	60.48	9.54	28.65	59.71	21.91	42.67	30.37	51.95	9.53	10.05	1.50	23.72	10.88	42.76	30.54	38.72	7.58	10.31	30.81	1.00	1.93	14.31	10.12	47.06	38.83	45.91	45.44	34.15	40.61	
4.46	2.29	2.87	3.61	2:32	2.27	1.76	0. <b>4</b> .0	0.95	8.09	0.87	1.00	4.98	3.82	2.34	0.83	39.83	37.52	4.40	0.93	1.29	1.00	1.10	2.79	2.35	7.38	69.6	41.41	3.39	2.02	4.49	1.21	1.87	1.97	2.37	
25.24	25.21	25.10	25.09	24.95	24.86	24.50	24.25	24.24	23.75	23.68	23.00	22.37	22.30	21.79	21.71	21.70	21.44	20.97	20.39	20.04	19.97	19.61	19.50	19.47	19.37	19.26	18.93	18.85	18.21	18.01	17.78	17.62	17.44	17.25	9
0.25	0.25	0.25	0.17	0.25	24.91	0.24	26.54	7.41	0.84	3.51	0.23	0.16	3.74	0.34	3.65	0.14	40.77	19.16	6.19	3.94	0.20	3.26	0.20	3.30	0.19	0.19	21.01	0.17	0.74	0.35	3.45	0.54	0.17	0.44	
0.01	0.01	0.0	0.0	0.0	8	10.0	60	33	5 6	0.15	0.0	0.0	0.17	0.02	0.17	0.01	1.90	0.91	0.30	0.20	0.01	0.17	0.0	0.17	0.01	0.0	=	0.0	20	0.02	0.19	0.03	0.01	0.03	
AA010383	AA070437	R15813	AA236986	W94486	H19826	Haffa7	T57803	200004	CA444	A4504625	T201498	AA630328	R44936	AA161097	A 4 40 1 26 1	H10072	T61938	#062EN	AA255954	R95684	AA461508	AA520979	D86970	H28091	A 4 495022	A 4 4 5 6 8 2 8	765736	R54492	AA620747	AA057195	AA487505	R60020	AA001376	AA489662	
430255	530958	53315	687972	357884	172517	251806	80843	2000	20000	2000	81050	855910	24204	591055	02070	48843	28000	259884	686552	199337	795831	82626	407264	467240	759163	800683	80338	39722	1049284	489078	839048	7.79b7	361899	823727	:
168	9 04	5 5	<u> </u>	: :	<u> </u>	? ?	• · ·	0/1	2 !	<u> </u>	2 7		5 4	5 5	7 6	184	ž ž	3 4	187	188	180	3 5	2	<u>.</u> 5	<u> </u>	3 3	ž á	80	107	9 0	9 9	3 5	50.	202	ļ ;

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

2	2	7	2	7	2	7	2	2	2	7	2	7	2	7	7	7	2	2	2	2	7	7	7	7	7	7	2	7	7	2	2	7	7	2	
5.21	12.21	34.18	7.22	5.65	69.9	28.36	25.15	19.22	15.13	18.34	6.87	15.54	23.25	28.45	28.16	5.10	27.66	4.48	12.18	18.27	29.19	18.18	7.18	26.50	7.37	25.70	28.39	2.23	3.21	5.10	0.58	7.04	26.39	20.98	
42.11	37.48	1.51	39.78	40.90	36.93	16.52	21.09	22.78	28.43	26.10	0.27	3.02	18.93	4.19	9.24	36.28	2.5	18.97	26.44	21.58	8.82	18.08	29.16	1.54	0.76	12.28	8.17	29.01	28.51	31.49	14.55	29.00	0.39	14.19	
4.26	1.87	14.65	2.86	2.84	3.94	2.63	0.69	4.69	1.53	0.42	36.64	24.69	1.00	10.07	5.00	0.88	2.99	17.85	2.42	. 0.64	2.08	3.39	3.08	10.97	30.80	0.65	1.86	7.14	6.31	1.33	22.57	1.18	10.40	1.96	
17.19	17.19	16.78	16.62	16.47	15.85	15.84	15.64	15.57	15.03	14.95	14.60	14.41	14.40	14.24	14.13	14.09	13.89	13.76	13.68	13.50	13.36	13.22	13.14	13.00	12.97	12.88	12.81	12.79	12.67	12.64	12.56	12.41	12.39	12.38	٢
0.86	1.06	0.11	0.29	1.14	0.40	2.94	5.98	0.30	3.91	3.94	0.54	1.64	41.0	3.51	17.58	2.19	4.17	0.13	1.31	3.95	8.41	0.26	1.05	90.0	0.36	0.13	8.32	0.71	0.25	1.99	0.13	4.99	9.44	0.12	
0.05	90.0	0.01	0.02	0.07	9.03	0.19	0.38	0.02	0.26	0.26	9.0	0.11	0.01	0.25	1.24	0.16	0.30	0.01	0.10	0.29	0.63	0.02	90.0	0.01	0.03	0.01	0.65	90.0	0.02	0.16	0.01	0.40	0.76	0.01	
AA609304	AA857163	AA481481	N26559	N51297	AA410437	AA504838	AA417713	AA045175	AA463445	H18932	AA035745	AA476543	AA434388	AA883800	AA774665	AA699601	W90705	R56149	R37738	AA427719	H15549	AA491227	AA883181	N52193	AA035730	AA017379	AA911236	R43915	AA621355	AA070435	AA453616	AA437212	AA620697	AA022496	
1031568	1410444	756629	266366	283070	753411	825726	746232	487766	811766	51599	364865	785293	770860	1461609	856289	436402	418004	41406	26806	771004	49499	824074	1467166	284288	364844	361122	1526789	33066	1065121	530954	795446	757428	1049185	364352	
203	707	205	206	207	208	508	210	211	212	213	214	215	216	217	218	219	220	221	222	223	224	225	226	227	228	229	230	231	232	233	234	235	236	237	

Table 2-11; Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line; Genes upregulated 5 fold or more in at least 2 cancer cell lines.

N N	1 71	. 7	. 7	C4	7	8	. 2	. ~		,	, ,	4 6	4 (	<b>v</b> (	4 (	ν (	N (	71	2	7	7	7	8	7	~	. 6			7 (	7	7	7	~	2	7		
11.09	17.55	23.47	20.10	12.88	16.24	19.11	14.01	12.09	7.03		20.33	74.11	16.34 24.04	7.83	13.60	4.94	16.62	10.20	12.02	20.10	8.36	5.22	21.77	5.14	E 78	6.7.5	50.00 51.00	53.70	15.64	0.48	1.65	17.42	21.42	5.78	1.12	!	
23.87	29.43	10.47	5.4.0	06.FI	2,00	7	13.10	, 50.04 , 60.04	18.51	20.61	10.18	5.70	16.85	22.78	17.96	9.89	15.37	20.21	15.78	10.22	0.1	22.31	000	6.53	3 6	21.63	9:4	6.31	14.23	7.21	17.15	11.85	90.9	20.48	20.18	2	
211	6.29	3.28	8.77	9.5	4	2.08	3.09	0.58	3.60	10.87	3.65	4.58	0.33	2.58	0.92	17.77	0.51	2.09	4.64	2.05	22.91		7	4	06.12	2.67	0.88	0.86	0.77	22.87	11.51	0.93	267	72 6	7.0	0.0	
12.36	12.24	12.10	12.06	12.02	11.98	11.82	11.80	11.71	11.67	11.63	11.61	11.46	11.17	11.06	10.90	10.87	10.84	10.83	10.81	10.79	40.75	10.70	10.66	10.50	10.45	10.36	10.32	10.31	10.21	10.19	10.11	10.07	40 C	2 5	10.00	) n. c	∞
3.72	0.12	0.31	16.58	1.69	16.82	23.53	3.89	0.18	11.24	0.20	27.64	0.11	0.34	0.95	3.90	5.68	6.12	0.31	9	9.0	3.32	5	5.93	8.09	12.13	0.44	0.87	4.21	5.55	9.47	0.12	, ,,	50.3	19.00	0.37	2.42	
0:30	0.01	0.03	1.38	0.14	1.40	1.99	0.33	0.02	96.0	0.02	2.38	0.01	0.03	0.09	0.36	0.52	7.20	5 6	3 6	0.0	0.86	0.01	0.56	0.77	1.16	9.0	90.0	0.41	, O	683	3 6	5 6	07.0	1.60	<u>\$</u>	0.24	
H10713	157927	AA778717	AA476576	R92201	AA706901	AA477227	H09940	AA161188	R44816	W93523	AA026582	AA179161	AA160670	AA427737	UARDZO	010041	4440434	\$ \$ 1 ! ! !	N74131	AA461513	N30751	AA680136	AA417920	AA489073	AA454949	N93721	AA281784	H54451	NA9389	4 6 4 4 7 0 4 0	8000000	AA158035	R94504	AA478036	AA598787	AA626167	
47264	71671	1049079	785368	195845	451649	739450	46461	591157	33839	357138	366971	609228	50773	770854	403333	CCCCE1	130/40	133420	238417	795840	257955	433155	767403	824802	814769	307138	712401	203184	277624	120002	55827	290040	197651	740457	898073	745248	
82	83	240	241	242	243	244	245	246	247	248	249	250	, t	25.2	1 2	2 2	<b>5</b>	CC.7	256	257	258	259	260	261	762	263	784	265	3 8	8 5	9	268	503	270	271	272	

Table 2-11; Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

7	2	7	7	7	7	7	2	7	7	7	2	7	7	2	7	7	7	7	7	7	2	7	7	7	7	2	7	7	8	7	2	7	7	7	
16.30	14.06	15.31	10.88	1.81	0.46	16.03	5.79	6.87	18.56	14.13	6.28	18.44	9.58	10.89	15.44	16.97	18.79	15.70	18.46	21.00	18.37	15.08	8.67	14,44	9.79	17.77	4.33	5.45	11.30	4.46	13.51	13.08	14.35	21.12	
12.15	3.07	13.69	17.62	19.71	23.34	9.94	22.17	18.90	9.16	9.81	21.91	6.58	17.50	15.63	11.85	1.00	5.97	7.97	1.04	5:32	1.12	1.5¢	15.51	12.53	16.72	7.70	10.86	19.73	11.63	14.83	13.04	10.73	0.36	1.00	
1.40	12.71	0.67	1.10	8.07	5.59	3.40	1.35	3.45	1.31	4.99	0.65	3.83	1.61	2.05	1.15	10.45	3.48	4.56	8.70	1.84	8.59	11.43	3.79	76.0	1.35	2.30	12.49	2.37	4.58	8.17	0.81	3.46	12.50	5.06	
9.95	9.95	68 6	9.87	98.6	9.80	9.79	77.6	9.74	89.6	9.64	9.62	9.62	9.56	9.53	9.48	9.47	0.41	9.41	9.40	9.39	9.36	9.34	9.32	9.31	9.29	9.26	9.22	9.18	9.17	9.15	9.12	9.09	9.07	9.07	თ
2.29	1.05	2.	3.82	0.09	0.21	0.57	0.21	12.26	5.97	0.17	6.12	0.10	0.16	0.77	1.40	0.09	7.14	5.85	0.09	5.02	0.08	90:0	0.19	0.22	6.35	4.10	0.97	0.87	9.89	3.21	2.44	0.26	0.26	0.09	
0.23	0.11	0.50	0.39	0.01	0.02	90.0	0.02	1.28	0.62	0.02	9.0	0.01	0.02	90.0	0.15	0.01	0.76	0.62	0.01	0.53	0.01	0.01	0.02	0.02	0.68	4.0	0.10	0.10	90.	0.35	0.27	0.03	0.03	0.01	
AA664007	AA625979	AA292226	N21633	R55220	AA464615	A1003636	AA120779	AA682533	AA620528	AA157261	AA133590	AA158342	R89490	R46000	AA219060	W16659	AA678348	AA204830	AA679180	AA136664	AA699926	AA284072	W74377	N50962	AA455242	A1017507	AA705047	R95962	AA446462	H10192	R96579	R69584	W44766	AA100698	
855391	745490	725877	266135	154720	812994	391987	490387	431242	951305	230500	512116	590727	195695	35804	630013	320495	430973	645565	866702	490959	435330	700792	346552	281162	814791	1627705	461488	230882	781047	46694	199577	141726	320865	490556	
273	274	275	276	112	278	279	280	281	282	283	284	285	286	287	288	289	280	291	292	293	38	295	298	297	208	583	000	Š	305	8	Š	305	306	307	

Table 2-11; Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

71	7	2	7	7	7	24	7	7	7	7	7	61	7	7	7	7	7	7	74	2	7	7	7	2	7	7	7	7	7	7	64	7	7	7	
14.67	5.38	9.46	1.10	16.60	5.40	11.94	8.45	14.43	8.30	18.10	15.48	7.50	14.35	18.86	3.90	17.99	11.27	12.36	17.21	6.37	14.40	16.24	19.27	2.96	13.09	11.26	18.07	17.7	14.93	15.23	9.48	6.36	13.11	13.46	
10.87	20.24	15.22	13.45	0:30	20.84	2.96	17.16	2.08	17.44	6.64	6.65	16.58	1.00	0.52	14.21	6.50	12.46	11.51	6.16	13.73	6.65	6.27	0.41	13.18	1.67	11.14	0.58	13.64	5.10	3.98	13.54	18.34	7.53	8.34	
1.65	1.54	24.	12.54	10.05	0.65	11.72	0.92	9.82	0.36	1.3	3.86	1.87	10.33	6.22	7.42	0.78	1.45	1.30	1.81	4.95	3.89	2.41	5.22	8.74	9.98	2.23	5.83	3.04	4.35	5.06	1.22	0.38	3.42	2.25	
9.06	9.05	9.04	9.03	8.98	8.98	8.87	8.84	8.81	8.70	8.69	8.66	8.65	8.56	8.53	8.51	8.42	8.39	8.39	8.39	8.35	8.32	8.31	8.30	8.29	8.25	8.21	8.15	8.13	8.13	8.09	80'8	8.03	8.02	8.02	10
3.21	3.62	10.62	11.71	17.81	0.21	5.97	3.32	2.16	39.21	0.80	48.92	34.28	0.09	5.11	0.14	6.98	9.51	3.36	0.08	0.19	4.39	13.45	10.75	1.09	0.15	1.49	0.14	0.25	21.34	10.44	6.11	0.22	0.28	90.0	
0.35	0.40	1.17	1.30	1.98	0.02	0.67	0.38	0.25	4.51	0.09	5.65	3.96	0.01	0.60	0.02	0.83	1.13	0.40	0.01	0.05	0.53	1.62	1.30	0.13	0.02	0.18	0.02	0.03	2.63	1.29	0.76	0.03	0.03	0.01	
N26515	AA192527	AA438175	AA172056	AA620485	AA416970	W68220	H41203	H16772	AA629999	AA778648	H95638	AA775863	AA496998	AA682321	T52311	N80764	W47364	AA504265	AA036974	AA485265	AA455786	N74617	AA488324	R61372	N50806	AA058576	AA131239	W96216	R41329	H54752	AA699427	AA457576	AA705825	H94944	
266259	627541	877632	594758	951241	730036	342640	175533	50227	884511	1048995	242952	878544	897546	462926	71977	300615	324717	825416	484535	840150	809557	296429	842968	37980	280882	380884	503581	358599	30148	203275	433253	838776	1292182	230261	
308	309	310	311	312	313	314	315	316	317	318	319	320	321	322	323	324	325	326	327	328	329	330	331	332	333	334	335	336	337	338	339	340	341	342	

Table 2-11; Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

7	7	2	7	7	7	7	7	7	7	7	7	7	7	7	7	2	73	2	7	2	2	7	7	2	7	2	7	7	7	7	7	2	7	7	
8.40	8.68	6.01	6.49	10.12	4.46	15.20	10.00	11.39	10.84	9.28	13.28	1.00	11.71	9.03	6.84	2.00	5.66	9.32	15.79	7.15	15.99	9.88	8.23	15.83	9.18	5.28	12.04	9.29	4.93	12.46	12.18	8.02	14.40	10.67	
14.68	13.94	15.21	11.97	9.24	7.24	7.43	8.93	4.17	99.2	12.63	3.58	14.11	8.00	11.89	14.72	8.59	16.02	12.70	5.74	4.98	1.00	12.36	10.88	5.39	0.92	4.41	8.97	11.96	6.94	4.68	<b>6</b> .84	13.66	0.63	0.08	
0.92	1.05	2.27	4.86	3.94	11.54	0.57	4.23	7.55	4.60	1.18	6.23	7.95	3.35	2.09	1.41	12.36	1.23	0.83	1.27	10.53	5.64	0.35	3.47	1.25	12.26	12.64	1.30	6.99	10.35	5.03	3.00	0.34	6.83	11.07	
8.00	7.89	7 83	77.7	77.7	7.75	7.73	7.72	7.70	7.70	7.70	7.70	7.69	7.68	7.67	7.66	7.65	7.64	7.61	7.60	7.55	7.54	7.53	7.53	7.49	7.46	7.4	7.44	7.41	7.41	7.39	7.34	7.34	7.29	7.27	1
8.81	2.29	5.20	2.89	09.0	5.81	0.14	0.35	24.59	13.42	2.05	1.68	0.08	261.61	0.53	0.25	1.03	18.06	0.08	5.46	9.	90.0	7.28	5.64	1.18	0.08	0.20	6.14	0.31	21.88	23.90	57.27	9.91	28.27	15.11	
1.10	0.29	99.0	0.37	0.08	0.75	0.02	0.05	3.19	1.74	0.27	0.22	0.01	8	0.07	0.03	0.13	2.36	0.01	0.72	0.01	0.01	0.97	0.75	0.16	0.01	0.03	0.83	9.	2.95	3.23	7.80	1.35	3.88	2.08	
H51765	AA284172	R50775	H29897	AA625764	AA431741	AA156795	AA906257	T96718	155607	H57082	W15305	AA463982	AA676460	H10231	H10068	AA780365	W72666	T95804	AA291773	AA070331	AA702802	R53455	AA434403	AA633836	AA485749	T61122	R12473	R49644	AA701455	N93924	AA488367	N62936	T66816	AA706968	
193892	701625	38833	190325	745393	782259	502444	1505919	121251	73596	204644	322461	810661	882510	46927	46740	1035664	345743	120695	725340	530820	448032	39833	770879	868169	811168	83686	128243	37538	435076	309288	843163	278729	66317	451907	
55	35	3	346	<del>-</del>	, E	2	350	38.	352	353	354	355	356	357	358	359	360	38.	362	363	364	365	8	367	368	389	370	371	372	373	374	375	376	377	

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Table 2-11; Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line; Genes upregulated 5 fold or more in at least 2 cancer cell lines.

8	7	7	7	7	7	2	7	7	7	2	7	2	73	2	7	7	7	7	7	7	7	7	7	7	7	7	7	7	7	7	7	7	7	8	
10.22	12.41	6.82	6.11	10.54	6.43	5.38	8.99	11.27	10.38	10.69	11.80	7.84	7.62	5.32	8.84	6.81	3.65	6.50	11.87	9.20	14.49	11.35	9.27	10.38	1.16	3.14	9.86	5.69	8.65	12.53	8.19	9.34	11.61	5.09	
9.92	2.06	13.06	12.65	7.03	13.46	11.90	9.60	7.61	8.89	6.35	8.75	0.03	11.28	2.34	11.35	12.50	5.87	9.95	7.50	7.07	2.67	6.52	9.22	8.38	5.14	11.26	4.41	10.31	9.72	1.10	1.51	8.48	7.15	13.88	
1.57	7.23	1.81	2.92	4.11	1.78	4.39	3.01	2.72	2.25	4.40	0.87	13.51	2.47	13.68	1.14	2.00	11.72	4.65	1.68	4.74	0.85	3.01	2.37	2.09	14.50	6.40	6.49	4.76	2.34	7.04	10.95	2.73	1.77	1.52	
7.24	7.23	7.23	7,23	7.23	7.22	7.22	7.20	7.20	71.17	7.15	7.14	7.13	7.12	7.11	7.11	7.11	7.08	7.03	7.02	7.00	7.00	96.9	6.95	6.95	6.93	6.93	6.92	6.92	6.90	6.89	6.88	6.85	6.84	6.83	12
0.35	2.20	1.17	8.99	5.80	0.31	33.83	6.75	5 4	15.96	0.10	0.11	105.24	14.44	0.62	1.51	95.95	14.78	90:0	2.01	100.32	0.28	44.0	0.68	36.26	0.07	0.39	3.69	7.46	3.78	90.0	2.74	4.75	0.83	0.25	
0.05	0.30	0.16	1 24	9	3 2	¥ 68	3 8	<b>t</b> 8	,	3 6	200	14.77	2.03	60.0	0.21	13.50	2.09	0.01	0.29	14.33	9.0	0:00	0.10	5.22	0.01	90.0	0.53	1.08	0.55	0.01	0.40	0.69	0.12	9.0	
N22766	R43308	H40224	A 4 2 5 8 0 0	T82072	02012	2011102	13/488	A4460952	AA433321	10240CAA	A A 677655	44804567	AA455237	AA449118	R96520	N66158	N45236	R74203	N53940	AI015359	W72556	W86196	T68878	AA485453	T58145	R56046	N20305	AA465238	AA886739	AA669222	AA452278	AA129135	A1025259	AA398406	
266631	32766	2007	F0600	75 A 20	340777	333144	705157	# 10A	070500	707070	807274	1416782	814779	785845	199644	278531	283173	143365	247050	1637343	345330	415908	82215	811067	79254	40773	264502	814225	1500162	856585	787938	58650	1641367	726791	
378	370	n Cor	8	- 6 6	383	8 8	g g	280	600	700	000	600	39.	363	60   60   60	ğ	395	396	397	308	968	400	401	405	403	\$ 2	405	406	407	408	409	410	411	412	•

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

7	7	2	7	~	2	7	2	8	7	7	7	2	7	7	2	7	2	7	2	7	7	7	7	2	7	۲3	7	2	7	2	2	7	7	2	
12.23	6.48	5.38	6.30	49.4	10.97	0.98	8.83	7.65	4.14	7.33	6.95	8.44	7.62	8.09	8.12	1.85	9.54	4.80	9.68	3.05	4.63	5.67	6.74	5.55	6.04	10.01	7.59	7.97	8.78	9.48	6.08	9.70	9.25	1.28	
5.03	12.71	10.24	4.78	9.34	3.78	7.25	1.82	4.68	8.01	10.73	9.66	4.39	4.47	0.58	3.71	10.05	9.16	4.7	7.59	5.69	69.6	8.97	11.33	11.59	2.65	5.16	9.29	8.32	9.13	5.78	11.14	6.70	4.97	12.25	
3.20	1.22	4.75	9.29	6.37	5.57	12.08	9.56	7.84	8.02	2.08	3.37	7.13	7.85	11.27	8.09	7.97	1.16	7.52	2.44	10.89	5.27	4.93	1.47	2.35	10.79	4.26	2.50	3.07	1.38	4.03	2.05	2.85	5.00	5.69	
6.82	6.80	6.79	6.79	6.79	6.77	6.77	6.74	6.72	6.72	6.71	6.66	6.65	6.65	6.64	6.64	6.62	6.62	6.59	6.57	6.54	6.53	6.52	6.51	6.50	6.49	6.48	6.46	6.45	6.43	6.43	6.42	6.41	6.41	6.41	13
8.14	2.79	0.08	0.34	8.93	27.70	1.65	10.08	0.30	3.61	44.7	65.62	18.79	4.67	7.75	4.52	0.07	63.60	0.07	0.12	0.42	7.31	5.64	0.24	6.55	8.71	8.44	0.12	7.26	27.63	1.70	0.45	6.74	0.82	0.21	
1.19	0.41	0.01	0.05	1.32	4.09	0.24	1.50	9.	95.0	1.1	9.85	2.82	0.70	1.17	0.68	0.01	9.61	0.01	0.02	90.0	1.12	0.41	9.0	9.	1.3	1.30	0.02	1.13	4.30	0.26	0.07	1.05	0.13	0.03	
R23055	H56903	AA256231	H71230	H08595	AA010065	AA454562	AA454597	H17139	AA609723	AA398949	AA465593	AA599092	AA456895	AA424790	H59203	AA448400	T55728	AA884622	AA039851	H05645	H20847	N48075	T51617	R43328	AA953357	H21943	AA428186	AA045508	AA626788	AA034115	AA452542	AA448664	AA757564	H15910	
131316	204580	681906	214604	46091	359119	809503	811582	51083	1031885	725677	814246	950445	813149	768172	204214	781362	73638	1468597	375827	43662	51460	281733	72441	32134	1573251	160233	773564	487297	877784	430007	788488	786078	395955	159462	
413	414	415	416	417	418	419	420	424	422	423	424	425	426	427	428	429	<b>63</b>	431	432	433	¥.	435	436	437	438	439	4	<b>1</b>	442	<b>44</b> 3	<b>‡</b>	445	84	447	

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

MAGNITGS         0.76         4.82         6.33         8.52         6.40         4.22         2           NUSSELI         0.59         3.76         6.37         1.54         6.37         1.54         6.37         1.54         6.37         1.54         6.37         1.54         6.38         1.54         6.38         1.204         2           AAAGSS3         0.27         1.66         6.36         0.76         1.520         6.13         2           AAAGSS44         0.77         4.87         6.31         1.06         8.03         8.86         2           AAAGS244         0.77         4.87         6.31         1.06         8.03         8.86         2           AAAGS244         0.77         4.87         6.31         1.06         8.03         8.86         2           AAAGS24         0.77         4.87         6.39         1.79         10.33         8.86         2           AAAGS24         0.77         4.87         6.39         1.74         10.13         2.32         8.63         8.63         8.86         2           NAGS24         0.72         6.23         1.74         10.13         7.26         8.89         2     <	53         0.76         4.82         6.38         6.52         6.40           13         0.59         3.76         6.37         2.87         7.88           33         0.71         4.54         6.37         1.54         5.53           44         0.72         4.54         6.36         0.76         1220           24         0.77         4.87         6.36         1.48         6.38           4         0.77         4.87         6.39         1.79         1.220           2         8.25         5.186         6.30         1.79         1.078           2         8.25         5.186         6.30         1.79         1.078           2         8.25         5.196         6.30         1.79         1.078           0.02         0.13         6.29         2.64         6.05         6.27         2.46         6.05           0.02         0.13         6.27         2.11         8.82         6.27         2.46         6.05           0.02         0.14         6.23         0.27         2.46         6.05         6.25         3.19         6.05           0.22         1.36         6.25         0.21 </th <th>194717</th> <th>R89846</th> <th>0.26</th> <th>1.66</th> <th>6.39</th> <th>06.0</th> <th>90.9</th> <th>12.22</th> <th>7</th>	194717	R89846	0.26	1.66	6.39	06.0	90.9	12.22	7
33         376         6.37         2.87         7.88           34         6.37         1.54         6.53         7.88           35         1.01         6.46         6.36         0.76         1.50         6.53           4         1.50         6.36         0.76         1.220         1.220           4         0.77         4.87         6.31         1.06         8.03           4         0.77         4.87         6.31         1.06         8.03           5         8.25         51.96         6.30         1.79         10.78           0.72         4.53         6.29         3.26         8.03           0.02         0.15         6.29         3.26         8.03           0.02         1.37         6.27         2.14         8.05           0.02         0.15         6.27         2.11         8.23           0.02         0.15         6.27         2.11         8.23           0.02         0.15         6.27         2.11         8.23           0.02         0.15         6.25         3.19         6.21           0.02         0.14         6.23         0.24         10.34	13         0.59         3.76         6.37         2.87         7.88           13         0.71         4.54         6.37         1.54         5.53           10         0.24         1.50         6.36         0.76         1.20           12         1.01         6.46         6.36         1.48         6.38           4         0.77         4.87         6.31         1.06         8.03           2         8.25         51.96         6.30         1.79         10.78           2         8.25         51.96         6.29         3.26         6.38           0.72         4.53         6.29         3.26         6.93           0.02         0.13         6.29         3.26         6.93           0.03         5.87         6.29         3.26         6.93           0.02         0.13         6.27         2.46         6.03           0.02         0.15         6.27         2.46         6.03           0.02         0.15         6.25         2.11         8.25           0.02         0.15         6.25         0.27         2.46           0.22         1.36         6.25         3.19		AA431753	0.76	4.82	6.38	8.52	6.40	4.22	7
13         0,71         4,54         6,37         1,54         5,53           10         0,24         1,50         6,36         0,76         12,20           4         0,77         4,87         6,36         1,48         6,38           4         0,77         4,87         6,31         1,06         6,38           5         8,25         51,96         6,30         1,79         10,78           0,72         4,53         6,29         3,26         6,33           0,03         5,87         6,29         3,26         6,93           0,03         2,21         6,29         2,64         8,05           0,03         2,21         6,29         2,64         8,05           0,02         1,37         6,27         2,64         8,05           0,02         1,37         6,27         2,14         8,25           0,02         1,58         6,27         2,14         8,25           0,02         1,58         6,27         2,14         8,25           0,02         1,36         6,25         3,19         6,21           0,02         1,36         6,25         3,19         6,21	13         0.71         4.54         6.37         1.54         5.53           10         0.24         1.50         6.36         0.76         12.20           24         1.50         6.36         0.76         12.20           4         0.77         4.87         6.31         1.06         8.03           2         8.25         51.96         6.30         1.79         10.78           2         8.25         51.96         6.30         1.79         10.78           0.02         6.23         6.29         3.26         6.83           0.03         5.87         6.29         3.26         6.83           0.03         5.87         6.29         3.26         6.83           0.02         1.37         6.23         2.64         8.05           0.02         1.37         6.27         2.40         2.46         8.05           0.02         1.58         6.27         2.11         8.27         3.05         4.65           0.02         1.58         6.25         3.19         6.25         3.19         6.25         3.19         6.24         6.25         3.16         6.21         3.24         6.25		N39611	0.59	3.76	6.37	2.87	7.88	8.37	7
3.24         1.50         6.36         0.76         12.20           4.6         6.36         6.36         1.48         6.38           4.7         6.31         1.06         6.36         1.48         6.38           4.7         6.31         1.06         8.03         1.20         6.38           2.2         8.25         51.96         6.30         1.79         10.78         6.38           0.72         4.53         6.29         3.26         6.93         10.78         6.93           0.35         2.21         6.29         2.64         8.05         6.93           0.22         1.37         6.27         7.40         2.46         8.05           0.25         1.58         6.27         2.11         8.2         2.46         8.05           0.02         0.15         6.27         2.11         8.2         2.46         8.05           0.03         6.25         1.31         8.2         2.46         8.05           0.04         1.36         6.25         2.11         8.2         2.46         8.05           1.94         1.201         6.23         6.25         0.21         1.24         1.24 </th <th>3.24         1.50         6.36         0.76         12.20           3.2         1.01         6.46         6.36         1.48         6.38           4         0.77         4.87         6.31         1.06         8.03           2         8.25         51.96         6.30         1.79         10.78           0.72         4.53         6.29         3.26         8.03           0.93         5.87         6.29         2.64         8.05           0.35         2.21         6.28         1.45         10.73           0.22         1.37         6.27         2.64         8.05           0.22         1.37         6.27         2.46         8.05           0.22         1.58         6.27         2.11         8.2           0.34         2.29         6.25         0.27         2.46           0.35         6.25         0.27         2.05           0.36         4.11         6.23         0.27         5.01           0.38         2.36         6.25         0.27         5.01           1.94         12.01         6.25         0.3         1.20         0.76           1.94         12.0</th> <th></th> <th>AA405533</th> <th>0.71</th> <th>4.54</th> <th>6.37</th> <th><del>7.</del></th> <th>5.53</th> <th>12.04</th> <th>7</th>	3.24         1.50         6.36         0.76         12.20           3.2         1.01         6.46         6.36         1.48         6.38           4         0.77         4.87         6.31         1.06         8.03           2         8.25         51.96         6.30         1.79         10.78           0.72         4.53         6.29         3.26         8.03           0.93         5.87         6.29         2.64         8.05           0.35         2.21         6.28         1.45         10.73           0.22         1.37         6.27         2.64         8.05           0.22         1.37         6.27         2.46         8.05           0.22         1.58         6.27         2.11         8.2           0.34         2.29         6.25         0.27         2.46           0.35         6.25         0.27         2.05           0.36         4.11         6.23         0.27         5.01           0.38         2.36         6.25         0.27         5.01           1.94         12.01         6.25         0.3         1.20         0.76           1.94         12.0		AA405533	0.71	4.54	6.37	<del>7.</del>	5.53	12.04	7
32         1,01         6,46         6,36         1,48         6,38           44         0,77         4,87         6,31         1,06         8,03           25         8,25         51,96         6,39         1,79         10,78           0,72         4,53         6,29         3,26         6,93           0,25         2,21         6,29         2,64         8,05           0,22         1,37         6,29         2,64         8,05           0,22         1,37         6,27         7,40         2,46           0,22         1,37         6,27         7,40         2,46           0,22         1,37         6,27         7,40         2,46           0,25         1,58         6,27         2,11         8,82           0,37         2,29         6,27         2,11         8,82           0,37         2,29         6,25         0,27         5,01           0,39         2,35         6,25         0,37         5,01           1,94         12,01         6,23         0,87         12,34           0,22         1,36         6,25         0,87         1,23           1,94         1,	1.01         6.46         6.36         1.48         6.38           4         0.77         4.87         6.31         1.06         8.03           2         8.25         51.96         6.30         1.79         10.78           0.72         4.53         6.29         3.26         6.93           0.03         5.87         6.29         2.64         8.05           0.22         1.37         6.28         2.64         8.05           0.22         1.37         6.29         2.64         8.05           0.22         1.37         6.27         7.40         2.46           0.02         0.15         6.27         7.40         2.46           0.02         0.15         6.27         2.11         8.2           0.02         0.15         6.25         0.27         2.0           0.34         2.35         6.25         3.19         6.05           0.02         0.14         6.23         0.27         5.01           0.24         4.19         6.25         3.19         6.05           1.94         12.01         6.25         0.27         5.0           0.24         6.15         4.19 </th <th></th> <th>R93068</th> <th>0.24</th> <th>1.50</th> <th>6.36</th> <th>0.76</th> <th>12.20</th> <th>6.13</th> <th>~</th>		R93068	0.24	1.50	6.36	0.76	12.20	6.13	~
4         0.77         4.87         6.31         1.06         8.03           2         8.25         51.96         6.30         1.79         10.78           0.72         4.53         6.29         3.26         6.93           0.03         5.87         6.29         3.26         6.93           0.02         1.37         6.28         1.45         10.13           0.22         1.37         6.27         7.40         2.46           0.22         1.58         6.27         7.40         2.46           0.22         1.58         6.27         7.40         2.46           0.02         0.15         6.27         7.40         2.46           0.02         0.15         6.27         2.11         8.2           0.03         2.29         6.25         0.27         5.01           0.34         1.369         6.25         0.27         5.01           0.25         1.36         6.25         0.27         5.01           0.26         4.11         6.23         0.68         3.24         6.24           0.27         1.27         0.24         6.15         1.20         0.06           0.2	4         0.77         4.87         6.31         1.06         8.03           2         8.25         51.96         6.30         1.79         10.78           0.72         4.53         6.29         3.26         6.93           0.03         5.87         6.29         2.64         8.05           0.22         1.37         6.28         1.45         10.13           0.22         1.37         6.27         7.40         2.46           0.22         1.37         6.27         7.40         2.46           0.22         1.37         6.27         7.40         2.46           0.22         1.37         6.27         7.40         2.46           0.02         0.15         6.27         2.11         8.82           0.02         0.15         6.27         2.51         6.21         0.27           0.33         2.35         6.25         0.27         5.01         0.27         0.27           0.42         2.35         6.25         0.27         5.01         0.24         0.25         0.04         0.24         0.25           1.94         1.20         6.25         0.24         1.34         0.24         <		AA465223	1.01	6.46	6.36	1.48	6.38	11.23	7
2         8.25         51.96         6.30         1.79         10.78           0.72         4.53         6.29         3.26         6.93           0.93         5.87         6.29         3.26         6.93           0.35         2.21         6.29         2.64         8.05           0.22         1.37         6.27         7.40         2.46           0.25         1.58         6.27         7.40         2.46           0.25         1.58         6.27         7.40         2.46           0.02         1.15         6.27         2.11         8.26           0.02         0.15         6.27         2.11         8.26           0.33         2.35         6.25         0.27         5.01           0.24         4.11         6.23         0.27         5.05           0.25         1.36         6.25         0.27         5.05           0.27         1.36         6.25         0.27         5.01           0.27         1.21         6.25         0.27         5.01           0.28         1.201         6.25         0.28         0.76           0.42         12.17         6.19 <td< th=""><th>2         8.25         51.96         6.30         1.79         10.78           0.72         4.53         6.29         3.26         6.93           0.93         5.87         6.29         3.26         6.93           0.25         1.37         6.28         1.45         10.13           0.22         1.37         6.27         7.40         2.46           0.25         1.58         6.27         7.40         2.46           0.25         1.58         6.27         7.40         2.46           0.02         0.15         6.27         2.11         8.82           0.02         0.15         6.27         2.51         6.21           0.33         2.35         6.25         0.27         5.01           0.34         2.35         6.25         0.27         5.01           0.22         1.36         6.25         0.87         12.34           0.24         1.20         0.68         0.27         5.01           1.94         1.20         6.25         0.87         12.34           0.05         0.05         6.13         6.13         6.13         6.13           1.97         1.21         <t< th=""><th></th><th>AA459244</th><th>0.77</th><th>4.87</th><th>6.31</th><th>1.06</th><th>8.03</th><th>9.86</th><th>7</th></t<></th></td<>	2         8.25         51.96         6.30         1.79         10.78           0.72         4.53         6.29         3.26         6.93           0.93         5.87         6.29         3.26         6.93           0.25         1.37         6.28         1.45         10.13           0.22         1.37         6.27         7.40         2.46           0.25         1.58         6.27         7.40         2.46           0.25         1.58         6.27         7.40         2.46           0.02         0.15         6.27         2.11         8.82           0.02         0.15         6.27         2.51         6.21           0.33         2.35         6.25         0.27         5.01           0.34         2.35         6.25         0.27         5.01           0.22         1.36         6.25         0.87         12.34           0.24         1.20         0.68         0.27         5.01           1.94         1.20         6.25         0.87         12.34           0.05         0.05         6.13         6.13         6.13         6.13           1.97         1.21 <t< th=""><th></th><th>AA459244</th><th>0.77</th><th>4.87</th><th>6.31</th><th>1.06</th><th>8.03</th><th>9.86</th><th>7</th></t<>		AA459244	0.77	4.87	6.31	1.06	8.03	9.86	7
0.72         4.53         6.29         3.26         6.93           0.93         5.87         6.29         2.64         8.05           0.25         2.21         6.28         1.45         10.13           0.22         1.37         6.27         7.40         2.46           0.25         1.58         6.27         7.40         2.46           0.25         1.58         6.27         2.11         8.82           0.02         0.15         6.27         2.11         8.82           0.03         1.36         6.25         0.27         5.01           0.24         1.36         6.25         0.27         5.01           0.25         1.36         6.25         0.27         5.05           0.01         0.08         6.25         0.07         5.50           0.01         0.08         6.20         1.36         0.76           0.01         0.08         6.20         1.35         0.76           0.01         0.08         6.20         1.05         0.76           0.01         0.08         6.20         1.06         0.76           0.02         1.21         6.13         6.16 <td< th=""><th>0.72         4.53         6.29         3.26         6.93           0.93         5.87         6.29         2.64         8.05           0.35         2.21         6.28         1.45         10.13           0.22         1.37         6.27         7.40         2.46           0.25         1.58         6.27         7.40         2.46           0.02         0.15         6.27         2.11         8.82           0.02         0.15         6.27         2.11         8.82           0.02         0.15         6.27         2.51         6.21         2.46           0.33         2.35         6.25         0.27         5.01         2.27         6.25         13.18         0.27         5.01           0.24         4.11         6.25         13.18         6.25         13.18         6.25         10.20         0.27         5.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         <td< th=""><th></th><th>AA042812</th><th>8.25</th><th>51.96</th><th>6.30</th><th>1.79</th><th>10.78</th><th>6.32</th><th>7</th></td<></th></td<>	0.72         4.53         6.29         3.26         6.93           0.93         5.87         6.29         2.64         8.05           0.35         2.21         6.28         1.45         10.13           0.22         1.37         6.27         7.40         2.46           0.25         1.58         6.27         7.40         2.46           0.02         0.15         6.27         2.11         8.82           0.02         0.15         6.27         2.11         8.82           0.02         0.15         6.27         2.51         6.21         2.46           0.33         2.35         6.25         0.27         5.01         2.27         6.25         13.18         0.27         5.01           0.24         4.11         6.25         13.18         6.25         13.18         6.25         10.20         0.27         5.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05         6.05 <td< th=""><th></th><th>AA042812</th><th>8.25</th><th>51.96</th><th>6.30</th><th>1.79</th><th>10.78</th><th>6.32</th><th>7</th></td<>		AA042812	8.25	51.96	6.30	1.79	10.78	6.32	7
0.93         587         629         264         8.05           0.35         2.21         6.28         1.45         10.13           0.22         1.37         6.27         7.40         2.46           0.25         1.58         6.27         7.40         2.46           0.02         0.15         6.27         2.11         8.82           0.02         0.15         6.25         0.27         5.01           0.37         2.29         6.25         0.27         5.01           0.38         2.35         6.25         0.27         5.01           0.22         1.36         6.25         0.87         5.05           0.05         4.11         6.23         0.87         5.05           0.07         1.36         6.25         0.87         7.60           1.94         12.01         6.20         11.05         0.75         7.60           0.05         0.07         6.13         6.15         1.20         0.76         0.76           1.94         12.01         6.13         4.19         5.59         0.76         0.76           0.05         0.33         12.43         6.15         4.19	0.93         5.87         6.29         2.64         8.05           0.35         2.21         6.28         1.45         10.13           0.22         1.37         6.27         7.40         2.46           0.25         1.58         6.27         7.40         2.46           0.02         0.15         6.27         2.11         8.82           0.02         0.15         6.27         2.11         8.82           0.03         1.369         6.25         0.27         5.01           0.38         2.35         6.25         0.27         5.05           0.02         4.11         6.23         0.87         7.50           0.01         0.08         6.20         1.35         7.60           0.01         0.08         6.20         1.35         7.60           0.01         0.08         6.20         1.35         7.60           0.01         0.08         6.20         1.35         7.60           0.01         0.08         6.20         1.105         0.76           0.01         0.08         6.10         4.19         5.59           0.05         0.33         6.16         1.20         <		N63744	0.72	4.53	6.29	3.26	6.93	8.69	7
0.35         2.21         6.28         1.45         10.13           0.22         1.37         6.27         7.40         2.46           0.25         1.58         6.27         7.40         2.46           0.02         0.15         6.27         2.11         8.82           0.02         0.15         6.27         2.11         8.82           0.38         2.29         6.25         0.27         5.01           0.38         2.35         6.25         0.27         5.01           0.22         1.36         6.25         3.19         6.05           0.24         4.11         6.23         0.87         12.34           0.05         4.11         6.23         0.87         12.34           0.01         0.08         6.20         11.36         12.34           0.01         0.08         6.20         11.05         0.76           0.05         0.27         6.19         4.19         5.59           0.05         0.24         6.15         6.15         7.62         5.90           0.07         0.24         6.13         1.59         7.76         5.90           0.02         1.49	0.35         2.21         6.28         1.45         10.13           0.22         1.37         6.27         7.40         2.46           0.25         1.58         6.27         7.40         2.46           0.02         0.15         6.27         2.11         8.82           0.02         0.15         6.27         2.11         8.82           0.03         2.29         6.25         0.27         5.01           0.38         2.35         6.25         0.27         5.01           0.22         1.36         6.25         0.87         12.34           0.05         4.11         6.23         0.87         12.34           0.05         4.11         6.23         0.87         7.60           0.01         0.08         6.20         11.05         0.76           1.94         12.01         6.13         4.19         5.59           0.01         0.08         6.20         11.05         0.76           1.94         12.07         6.15         4.19         5.59           0.05         0.33         6.16         6.15         7.62         5.90           0.07         2.89         6.13		W85878	0.93	5.87	6.29	2.64	8.05	8.17	~
0.22         1.37         6.27         7.40         2.46           0.25         1.58         6.27         2.11         8.82           0.02         0.15         6.27         2.11         8.82           0.02         0.15         6.27         2.51         6.21           0.37         2.29         6.25         0.27         5.01           0.38         2.35         6.25         3.19         6.02           0.22         1.36         6.25         3.19         6.05           0.22         1.36         6.25         0.87         12.34           0.02         1.36         6.20         1.35         7.60           1.94         12.01         6.20         1.35         7.60           0.01         0.08         6.20         1.105         0.76           1.97         12.17         6.19         4.19         5.59           0.05         0.33         6.16         1.20         7.65           1         2.03         6.15         1.20         7.65           0.05         15.94         6.13         1.59         7.76           0.07         2.69         6.13         1.59 <t< td=""><td>0.22         1.37         6.27         7.40         2.46           0.25         1.58         6.27         2.11         8.82           0.02         0.15         6.27         2.11         8.82           0.02         0.15         6.27         2.51         6.21           0.37         2.29         6.25         0.27         5.01           0.38         2.35         6.25         3.19         6.02           0.38         2.35         6.25         3.19         6.05           0.22         1.36         6.25         0.87         12.34           0.02         1.36         6.20         1.35         7.60           0.01         0.08         6.20         1.35         7.60           1.94         12.17         6.19         4.19         5.59           0.01         0.08         6.20         11.05         0.76           1.97         12.17         6.19         4.19         5.59           0.05         0.33         6.16         6.15         7.62         5.90           0.47         2.89         6.13         1.59         7.76         7.62         5.90           0.02</td><th></th><th>N63988</th><td>0.35</td><td>2.21</td><td>6.28</td><td>1.45</td><td>10.13</td><td>7.26</td><td>2</td></t<>	0.22         1.37         6.27         7.40         2.46           0.25         1.58         6.27         2.11         8.82           0.02         0.15         6.27         2.11         8.82           0.02         0.15         6.27         2.51         6.21           0.37         2.29         6.25         0.27         5.01           0.38         2.35         6.25         3.19         6.02           0.38         2.35         6.25         3.19         6.05           0.22         1.36         6.25         0.87         12.34           0.02         1.36         6.20         1.35         7.60           0.01         0.08         6.20         1.35         7.60           1.94         12.17         6.19         4.19         5.59           0.01         0.08         6.20         11.05         0.76           1.97         12.17         6.19         4.19         5.59           0.05         0.33         6.16         6.15         7.62         5.90           0.47         2.89         6.13         1.59         7.76         7.62         5.90           0.02		N63988	0.35	2.21	6.28	1.45	10.13	7.26	2
0.25         1.58         6.27         2.11         8.82           0.02         0.15         6.27         2.51         6.21           0.03         2.29         6.25         0.27         5.01           1.219         13.69         6.25         13.18         6.27           0.28         2.35         6.25         3.19         6.02           0.22         1.36         6.23         0.87         12.34           0.22         1.36         6.22         0.68         8.92           1.94         12.01         6.20         11.35         0.76           0.01         0.08         6.20         11.35         0.76           0.02         12.01         6.19         4.19         5.59           0.04         12.07         6.15         4.19         5.59           0.05         0.33         6.16         1.20         1.069           0.42         2.57         6.15         4.49         5.59           0.42         2.57         6.15         4.26         5.90           0.02         0.14         6.13         1.59         7.76           0.03         2.40         6.11         2.39	0.25         1.58         6.27         2.11         882           0.02         0.15         6.27         2.51         6.21           0.03         2.29         6.25         0.27         5.01           0.38         2.35         6.25         3.19         6.05           0.05         4.11         6.23         0.87         12.34           0.02         1.36         6.22         0.68         8.92           1.94         12.01         6.20         1.35         7.60           0.01         0.08         6.20         1.35         7.60           0.05         0.33         6.16         4.19         5.59           0.05         0.33         6.16         1.20         7.65           1         2.60         1.20         4.19         5.59           0.05         0.33         6.16         4.19         5.59           0.42         2.57         6.15         4.19         5.59           0.42         2.57         6.15         4.19         5.59           0.42         2.57         6.15         7.62         5.90           0.42         2.54         6.15         7.62         5.		N77828	0.22	1.37	6.27	7.40	2.46	8.95	7
0.02         0.15         6.27         2.51         6.21           0.37         2.29         6.25         0.27         5.01           0.38         2.35         6.25         3.19         5.05           0.05         4.11         6.23         0.87         5.05           0.02         4.11         6.23         0.87         12.34           0.02         1.36         6.22         0.68         8.92           1.94         12.01         6.20         1.35         7.60           0.01         0.08         6.20         1.35         7.60           0.01         1.201         6.20         1.35         7.60           0.02         0.33         6.16         1.20         0.76           0.05         0.33         6.16         1.20         1.069           0.05         0.35         6.15         8.48         2.18           0.07         1.54         6.13         6.15         7.55           0.07         0.14         6.13         1.59         7.76           0.08         2.40         6.11         1.12         9.73           0.81         4.36         6.08         1.52	0.02         0.15         6.27         2.51         6.21           0.37         2.29         6.25         0.27         5.01           0.38         2.35         6.25         3.19         5.05           0.36         4.11         6.23         0.87         12.34           0.22         1.36         6.22         0.68         8.92           1.94         12.01         6.20         1.35         7.60           0.01         0.08         6.20         1.35         7.60           1.94         12.01         6.20         1.35         7.60           0.01         0.08         6.20         1.35         7.60           0.01         0.08         6.20         1.35         7.60           0.02         0.33         6.16         1.20         1.06           0.04         2.57         6.15         8.48         2.18           0.05         0.35         6.13         6.15         7.55           0.07         0.14         6.13         1.59         7.76           0.08         2.40         6.11         1.12         3.73           0.81         4.36         6.08         1.52         <		H54659	0.25	1.58	6.27	2.11	8.82	7.88	7
0.37         2.29         6.25         0.27         5.01           0.38         2.35         6.25         13.18         0.27           0.38         2.35         6.25         13.18         0.27           0.05         4.11         6.23         0.87         12.34           0.22         1.36         6.22         0.68         8.92           1.94         12.01         6.20         1.35         7.60           0.01         0.08         6.20         1.35         7.60           0.01         0.08         6.20         1.35         7.60           0.02         12.17         6.19         4.19         5.59           0.03         6.16         1.20         1.069         5.59           0.04         2.57         6.15         8.48         2.18           0.05         12.43         6.13         6.15         7.55           0.07         2.89         6.13         1.59         7.76           0.08         2.40         6.11         1.12         8.73           0.07         2.654         6.09         1.52         11.73           0.01         0.08         6.06         6.43	0.37         2.29         6.25         0.27         5.01           0.38         2.35         6.25         13.18         0.27           0.38         2.35         6.25         13.18         0.27           0.22         1.36         6.25         3.19         5.05           0.22         1.36         6.22         0.68         8.92           1.94         12.01         6.20         1.35         7.60           0.01         0.08         6.20         1.35         7.60           0.01         0.08         6.19         4.19         5.59           0.02         0.33         6.16         1.20         1.069           0.05         0.33         6.16         1.20         1.069           0.05         0.33         6.15         8.48         2.18           0.07         12.43         6.13         1.59         7.76           0.07         0.47         2.89         6.13         1.59         7.76           0.02         0.14         6.12         7.62         5.90           0.03         2.40         6.11         1.12         9.73           0.81         4.93         6.06		R44201	0.02	0.15	6.27	2.51	6.21	10.08	7
7         2.19         13.69         6.25         13.18         0.27           9.38         2.35         6.25         3.19         5.05           9 0.66         4.11         6.25         0.87         12.34           9 0.22         1.36         6.22         0.68         8.92           1 1.94         12.01         6.20         1.35         7.60           0 0.01         0.08         6.20         11.05         0.76           0 0.02         0.33         6.16         4.19         5.59           0 0.05         0.33         6.16         1.20         10.69           0 0.47         2.57         6.15         8.48         2.18           0 0.47         2.89         6.13         1.59         7.76           0 0.47         2.89         6.13         1.59         7.76           0 0.2         0.14         6.12         7.62         5.90           0 0.3         2.40         6.11         1.12         8.73           0 0.3         2.40         6.11         1.12         8.73           0 0.3         2.6.54         6.09         1.52         11.73           0 0.1         0.06	7         2.19         13.69         6.25         13.18         0.27           0.38         2.35         6.25         3.19         5.06           0.66         4.11         6.25         0.87         12.34           0.22         1.36         6.22         0.68         8.92           1.94         12.01         6.20         1.35         7.60           0.01         0.08         6.20         11.05         0.76           0.02         0.33         6.16         4.19         5.59           0.05         0.33         6.16         1.20         10.69           0.05         0.33         6.16         1.20         10.69           0.47         2.89         6.13         1.59         7.76           0.07         0.04         6.13         1.59         7.76           0.02         0.14         6.12         7.62         5.90           0.02         0.14         6.12         7.62         5.90           0.03         2.40         6.11         1.12         9.73           0.81         4.93         6.06         6.43         4.55           0.11         7.89         6.06		N24869	0.37	2.29	6.25	0.27	5.01	13.49	7
0.38         2.35         6.25         3.19         5.05           0.66         4.11         6.23         0.87         12.34           0.22         1.36         6.22         0.68         8.92           1.94         12.01         6.20         1.35         7.60           0.01         0.08         6.20         11.05         0.76           1.97         12.17         6.19         4.19         5.59           0.05         0.33         6.16         1.20         10.69           1         2.60         15.94         6.15         2.18           1         2.03         15.34         6.15         2.18           0.47         2.89         6.13         6.15         7.76           0.02         0.14         6.12         7.62         5.90           0.03         2.40         6.11         7.62         5.90           0.36         2.65         6.09         1.52         11.73           0.31         2.65         6.09         1.52         11.73           0.01         0.08         6.06         6.43         4.55           0.13         6.06         6.43         4.55	0.38         2.35         6.25         3.19         5.05           0.66         4.11         6.23         0.87         12.34           0.22         1.36         6.22         0.68         8.92           1.94         12.01         6.20         1.35         7.60           0.01         0.08         6.20         11.05         0.76           1.97         12.17         6.19         4.19         5.59           0.05         0.33         6.16         1.20         10.69           1         2.60         15.94         6.13         6.15         2.18           1         2.60         15.94         6.13         6.15         7.65           0.47         2.89         6.13         6.15         7.76           0.02         0.14         6.12         7.62         5.90           0.39         2.40         6.11         1.12         9.73           0.39         2.65         6.09         1.52         11.73           0.31         4.93         6.06         6.43         4.55           0.11         7.89         6.06         6.43         4.55           1.31         7.89 <t< td=""><th></th><th>AA430367</th><td>2.19</td><td>13.69</td><td>6.25</td><td>13.18</td><td>0.27</td><td>5.30</td><td>7</td></t<>		AA430367	2.19	13.69	6.25	13.18	0.27	5.30	7
9         0.66         4.11         6.23         0.87         12.34           0.22         1.36         6.22         0.68         8.92           1.94         12.01         6.20         1.35         7.60           0.01         0.08         6.20         11.05         0.76           1.97         12.17         6.19         4.19         5.59           0.05         0.33         6.16         1.20         10.69           1         2.03         6.15         8.48         2.18           1         2.03         6.15         8.48         2.18           0.47         2.89         6.13         0.57         7.55           0.02         0.14         6.13         0.57         7.56           0.03         2.40         6.13         1.59         7.76           0.36         2.40         6.11         1.12         9.73           0.35         2.65         6.09         1.52         11.73           0.81         4.93         6.06         6.43         4.55           0.11         7.89         6.06         6.43         4.55           0.16         0.94         6.02         3.5	9         0.66         4.11         6.23         0.87         12.34           0.22         1.36         6.22         0.68         8.92           1.94         12.01         6.20         1.35         7.60           0.01         0.08         6.20         11.05         0.76           0.05         0.33         6.16         4.19         5.59           0         0.42         2.57         6.15         8.48         2.18           1         2.60         15.94         6.13         6.57         7.55           1         2.03         12.43         6.13         6.15         2.69           0.047         2.89         6.13         6.15         7.76           0.02         0.14         6.12         7.62         5.90           0.03         2.40         6.11         1.12         9.73           2.62         16.02         6.11         1.12         9.73           0.39         2.40         6.11         1.12         9.73           0.81         4.93         6.06         6.43         4.55           0.11         0.09         6.06         6.43         4.56           0.16<		R11605	0.38	2.35	6.25	3.19	5.05	10.52	7
0.22         1.36         6.22         0.68         8.92           1.94         12.01         6.20         1.35         7.60           0.01         0.08         6.20         11.05         7.60           1.97         12.17         6.19         4.19         5.59           0.05         0.33         6.16         1.20         10.69           1         2.60         15.94         6.13         8.48         2.18           1         2.60         15.94         6.13         6.15         2.69           0.47         2.89         6.13         1.59         7.76           0.02         0.14         6.12         7.62         5.90           0.03         2.40         6.11         1.12         8.73           2.62         16.02         6.11         1.12         8.73           0.39         2.654         6.09         1.52         11.73           0.01         0.08         6.06         6.43         4.55           1.31         7.89         6.06         6.43         4.55           0.16         0.94         6.02         3.63         8.09	0.22         1.36         6.22         0.68         8.92           1.94         12.01         6.20         1.35         7.60           0.01         0.08         6.20         11.05         7.60           1.97         12.17         6.19         4.19         5.59           0.05         0.33         6.16         1.20         10.69           1         2.60         15.94         6.13         8.48         2.18           1         2.60         15.94         6.13         6.15         2.69           0.47         2.89         6.13         6.15         2.69           0.02         0.14         6.12         7.62         5.90           0.03         2.40         6.11         1.12         8.73           2.62         16.02         6.11         1.12         8.73           0.39         2.40         6.11         1.12         8.73           0.81         4.93         6.06         3.55         6.77           0.01         0.08         6.06         6.43         4.55           1.31         7.89         6.06         6.43         4.55           1.31         7.80 <td< td=""><th></th><th>AA701978</th><td>99.0</td><td>4.11</td><td>6.23</td><td>0.87</td><td>12.34</td><td>5.48</td><td>7</td></td<>		AA701978	99.0	4.11	6.23	0.87	12.34	5.48	7
1.94         12.01         6.20         1.35         7.60           0.01         0.08         6.20         11.05         0.76           0.05         0.33         6.16         4.19         5.59           1         0.42         2.57         6.15         8.48         2.18           1         2.60         15.94         6.13         6.15         7.55           1         2.03         12.43         6.13         6.15         2.69           0.047         2.89         6.13         1.59         7.76           0.02         0.14         6.12         7.62         5.90           0.03         2.40         6.11         1.12         9.73           2.62         16.02         6.11         1.12         9.73           0.39         2.40         6.11         1.12         9.73           0.81         4.93         6.06         1.52         11.73           0.01         0.08         6.06         6.43         4.55           0.15         7.89         6.06         6.43         4.55           0.16         0.94         6.02         3.03	1,94         12.01         6.20         1.35         7.60           0.01         0.08         6.20         11.05         0.76           0.05         0.33         6.16         4.19         5.59           0.05         0.33         6.15         8.48         2.18           1         2.60         15.94         6.13         0.57         7.55           1         2.60         15.94         6.13         6.15         2.69           0.47         2.89         6.13         1.59         7.76           0.02         0.14         6.12         7.62         5.90           0.03         2.40         6.11         1.12         9.73           2.62         16.02         6.11         1.12         9.73           0.39         2.40         6.11         1.12         9.73           0.30         2.40         6.11         1.12         9.73           0.81         4.93         6.06         1.52         11.73           0.01         0.08         6.06         6.43         4.55           1.31         7.89         6.04         4.66         8.09           0.16         0.94		R38539	0.22	1.36	6.22	0.68	8.92	9.06	8
0.01         0.08         6.20         11.05         0.76           1.97         12.17         6.19         4.19         5.59           0.05         0.33         6.16         1.20         10.69           1         0.42         2.57         6.15         8.48         2.18           1         2.60         15.34         6.13         0.57         7.55           1         2.03         12.43         6.13         6.15         2.69           0.07         0.14         6.12         7.62         5.90           0.02         0.14         6.12         7.62         5.90           0.39         2.40         6.11         1.12         9.73           2.62         16.02         6.11         1.12         9.73           0.39         2.6.54         6.09         1.52         11.73           0.01         0.08         6.06         6.43         4.55           1.31         7.89         6.04         4.66         8.09           0.16         0.94         6.02         3.62         3.03	0.01         0.08         6.20         11.05         0.76           1.97         12.17         6.19         4.19         5.59           0.05         0.33         6.16         1.20         10.69           1         0.42         2.57         6.15         8.48         2.18           1         2.60         15.94         6.13         0.57         7.55           1         2.03         12.43         6.13         6.15         2.69           0.07         0.07         6.13         1.59         7.76           0.08         6.13         1.59         7.76           0.09         6.11         1.12         9.73           0.09         6.11         1.12         9.73           0.39         2.40         6.11         1.12         9.73           0.36         6.05         6.11         2.39         8.86           0.36         6.06         6.15         1.52         11.73           0.81         4.93         6.06         6.43         4.55           0.13         7.89         6.06         6.43         4.55           0.16         0.94         6.02         9.62         3.0		H22568	<u>z</u> .	12.01	6.20	1.35	7.60	99.6	7
1,97         12,17         6,19         4,19         5.59           0,05         0,33         6,16         1,20         10,69           1         0,62         2,57         6,15         8,48         2,18           1         2,60         15,34         6,13         0,57         7,55           1         2,03         12,43         6,13         6,15         2,69           0,07         0,14         6,12         7,62         5,90           0,39         2,40         6,11         1,12         9,73           0,36         2,60         6,11         1,12         9,73           0,81         4,93         6,06         1,52         11,73           0,01         0,06         6,06         6,43         4,55           1,31         7,89         6,06         6,43         4,55           1,31         7,89         6,04         4,66         8,09           0,16         0,94         6,02         9,62         3,03	1,97         12,17         6,19         4,19         5.59           0,05         0,33         6,16         1,20         10,69           1         0,62         2,57         6,15         8,48         2,18           1         2,60         15,34         6,13         0,57         7,55           2         2,03         12,43         6,13         6,15         2,69           0,47         2,89         6,13         1,59         7,76           0,02         0,14         6,12         7,62         5,90           0,39         2,40         6,11         1,12         9,73           2,62         16,02         6,11         1,12         9,73           0,39         2,6,54         6,09         1,52         11,73           0,81         4,93         6,06         6,43         4,55           1,31         7,89         6,06         6,43         4,55           1,31         7,89         6,04         4,66         8,09           0,16         0,94         6,02         9,62         3,03		W31389	0.01	0.08	6.20	11.05	0.76	6.77	7
0.05         0.33         6.16         1.20         10.69           1         0.42         2.57         6.15         8.48         2.18           1         2.60         15.94         6.13         0.57         7.55           0.47         2.89         6.13         6.15         2.69           0.02         0.14         6.12         7.62         5.90           0.39         2.40         6.11         1.12         9.73           2.62         16.02         6.11         1.12         9.73           4.36         26.54         6.09         1.52         11.73           0.81         4.93         6.06         6.43         4.55           1.31         7.89         6.06         6.43         4.55           1.31         7.89         6.04         4.66         8.09           0.16         0.94         6.02         9.62         3.03	0.05         0.33         6.16         1.20         10.69           1         0.42         2.57         6.15         8.48         2.18           1         2.60         15.94         6.13         0.57         7.55           2.03         12.43         6.13         6.15         2.69           0.07         2.89         6.13         1.59         7.76           0.02         0.14         6.12         7.62         5.90           0.38         2.40         6.11         1.12         9.73           2.62         16.02         6.11         1.12         9.73           4.36         26.54         6.09         1.52         11.73           0.81         4.93         6.06         6.43         4.55           1.31         7.89         6.04         4.66         8.09           1.31         7.89         6.04         4.66         8.09           0.16         0.94         6.02         9.62         3.03		AA447662	1.97	12.17	6.19	4.19	5.59	8.79	7
0.42         2.57         6.15         8.48         2.18           2.60         15.34         6.13         0.57         7.55           1.203         12.43         6.13         0.57         7.55           0.47         2.89         6.13         1.59         7.76           0.02         0.14         6.12         7.62         5.90           0.39         2.40         6.11         1.12         9.73           2.62         16.02         6.11         1.12         9.73           4.36         26.54         6.09         1.52         11.73           0.81         4.93         6.06         6.43         4.55           0.01         0.08         6.06         6.43         4.55           1.31         7.89         6.04         4.66         8.09           0.16         0.94         6.02         9.62         3.03	0.42         2.57         6.15         8.48         2.18           2.60         15.94         6.13         0.57         7.55           1.203         12.43         6.13         0.57         7.55           0.47         2.89         6.13         1.59         7.76           0.02         0.14         6.12         7.62         5.90           0.39         2.40         6.11         1.12         9.73           2.62         16.02         6.11         1.12         9.73           4.36         26.54         6.09         1.52         11.73           0.81         4.93         6.06         6.43         4.55           0.01         0.08         6.06         6.43         4.55           1.31         7.89         6.04         4.66         8.09           0.16         0.94         6.02         9.62         3.03		N56875	9.05	0.33	6.16	120	10.69	<b>6</b> .60	7
2.60         15.94         6.13         0.57         7.55           2.03         12.43         6.13         6.15         2.69           0.047         2.89         6.13         1.59         7.76           0.02         0.14         6.12         7.62         5.90           0.39         2.40         6.11         1.12         9.73           2.62         16.02         6.11         1.12         9.73           4.36         26.54         6.09         1.52         11.73           0.81         4.93         6.06         6.43         4.55           0.01         0.08         6.06         6.43         4.55           1.31         7.89         6.04         4.66         8.09           0.16         0.94         6.02         9.62         3.03	2.60     15.94     6.13     0.57     7.55       2.03     12.43     6.13     6.15     2.69       0.07     2.89     6.13     1.59     7.76       0.02     0.14     6.12     7.62     5.90       0.39     2.40     6.11     1.12     9.73       2.62     16.02     6.11     2.39     8.86       4.36     26.54     6.09     1.52     11.73       0.01     0.08     6.06     6.43     4.55       1.31     7.89     6.04     4.66     8.09       0.16     0.94     6.02     9.62     3.03		AA865924	0.42	2.57	6.15	8.48	2.18	7.78	7
2.03         12.43         6.13         6.15         2.69           0.47         2.89         6.13         1.59         7.76           0.02         0.14         6.12         7.62         5.90           0.39         2.40         6.11         1.12         8.73           2.62         16.02         6.11         1.12         8.73           4.36         26.54         6.09         1.52         11.73           0.81         4.93         6.06         3.55         6.77           0.01         0.08         6.06         6.43         4.55           1.31         7.89         6.04         4.66         8.09           0.16         0.94         6.02         9.62         3.03	2.03     12.43     6.13     6.15     2.69       0.47     2.89     6.13     1.59     7.76       0.02     0.14     6.12     7.62     5.90       0.39     2.40     6.11     1.12     8.73       2.62     16.02     6.11     1.12     8.73       4.36     26.54     6.09     1.52     11.73       0.81     4.93     6.06     3.55     6.77       0.01     0.08     6.06     6.43     4.55       1.31     7.89     6.04     4.66     8.09       0.16     0.94     6.02     9.62     3.03       1.4		AA630104	2.60	15.94	6.13	0.57	7.55	10.29	2
0.47         2.89         6.13         1.59         7.76           0.02         0.14         6.12         7.62         5.90           0.39         2.40         6.11         1.12         9.73           2.62         16.02         6.11         2.39         8.86           4.36         26.54         6.09         1.52         11.73           0.81         4.93         6.06         6.43         4.55           0.01         0.08         6.06         6.43         4.55           1.31         7.89         6.04         4.66         8.09           0.16         0.94         6.02         9.62         3.03	0.47         2.89         6.13         1.59         7.76           0.02         0.14         6.12         7.62         5.90           0.39         2.40         6.11         1.12         8.73           2.62         16.02         6.11         2.39         8.86           4.36         26.54         6.09         1.52         11.73           0.81         4.93         6.06         6.43         4.55           0.01         0.08         6.06         6.43         4.55           1.31         7.89         6.04         4.66         8.09           0.16         0.94         6.02         9.62         3.03		AA430744	2.03	12.43	6.13	6.15	2.69	9.56	7
0.02     0.14     6.12     7.62     5.90       0.38     2.40     6.11     1.12     9.73       2.62     16.02     6.11     1.12     9.73       4.36     26.54     6.09     1.52     11.73       0.81     4.93     6.06     3.55     6.77       0.01     0.08     6.06     6.43     4.55       1.31     7.89     6.04     4.66     8.09       0.16     0.94     6.02     9.62     3.03	0.02     0.14     6.12     7.62     5.90       0.38     2.40     6.11     1.12     9.73       2.62     16.02     6.11     1.12     9.73       4.36     26.54     6.09     1.52     11.73       0.81     4.93     6.06     6.43     4.55       0.01     0.08     6.06     6.43     4.55       1.31     7.89     6.04     4.66     8.09       0.16     0.94     6.02     9.62     3.03		AA609891	0.47	2.89	6.13	1.59	7.76	9.04	7
0.39     2.40     6.11     1.12     9.73       2.62     16.02     6.11     1.12     9.73       4.36     26.54     6.09     1.52     11.73       0.81     4.93     6.06     3.55     6.77       0.01     0.08     6.06     6.43     4.55       1.31     7.89     6.04     4.66     8.09       0.16     0.94     6.02     9.62     3.03	0.39     2.40     6.11     1.12     9.73       2.62     16.02     6.11     1.12     9.73       4.36     26.54     6.09     1.52     11.73       0.81     4.93     6.06     3.55     6.77       0.01     0.08     6.06     6.43     4.55       1.31     7.89     6.04     4.66     8.09       0.16     0.94     6.02     9.62     3.03       1.4		H18017	0.02	0.14	6.12	7.62	5.90	4.83	7
2.62     16.02     6.11     2.39     8.86       4.36     26.54     6.09     1.52     11.73       0.81     4.93     6.05     3.55     6.77       0.01     0.08     6.06     6.43     4.55       1.31     7.89     6.04     4.66     8.09       0.16     0.94     6.02     9.62     3.03	2.62     16.02     6.11     2.39     8.86       4.36     26.54     6.09     1.52     11.73       0.81     4.93     6.05     3.55     6.77       0.01     0.08     6.06     6.43     4.55       1.31     7.89     6.04     4.66     8.09       0.16     0.94     6.02     9.62     3.03       14		R92186	0.39	2.40	6.11	1.12	9.73	7.48	2
4.36       26.54       6.09       1.52       11.73         0.81       4.93       6.06       3.55       6.77         0.01       0.08       6.06       6.43       4.55         1.31       7.89       6.04       4.66       8.09         0.16       0.94       6.02       9.62       3.03	4.36       26.54       6.09       1.52       11.73         0.81       4.93       6.06       3.55       6.77         0.01       0.08       6.06       6.43       4.55         1.31       7.89       6.04       4.66       8.09         0.16       0.94       6.02       9.62       3.03         1.4		AA430052	2.62	16.02	6.11	2.39	8.86	7.07	7
0.81     4.93     6.06     3.55     6.77       0.01     0.08     6.06     6.43     4.55       1.31     7.89     6.04     4.66     8.09       0.16     0.94     6.02     9.62     3.03	0.81     4.93     6.06     3.55     6.77       0.01     0.08     6.06     6.43     4.55       1.31     7.89     6.04     4.66     8.09       0.16     0.94     6.02     9.62     3.03       1.4		AA459106	4.36	26.54	6.09	1.52	11.73	5.03	7
0.01     0.08     6.06     6.43     4.55       1.31     7.89     6.04     4.66     8.09       0.16     0.94     6.02     9.62     3.03	0.01     0.08     6.06     6.43     4.55       1.31     7.89     6.04     4.66     8.09       0.16     0.94     6.02     9.62     3.03       14		H96392	0.81	4.93	90.9	3.55	6.77	7.86	7
1.31         7.89         6.04         4.66         8.09           0.16         0.94         6.02         9.62         3.03	1.31     7.89     6.04     4.66     8.09       0.16     0.94     6.02     9.62     3.03       14		AA102068	0.01	0.08	90'9	6.43	4.55	7.19	2
0.16 0.94 6.02 9.62 3.03	<b>0.16</b> 0.94 6.02 9. <b>62</b> 3.03		AA425404	1.31	7.89	6.04	4.66	8.09	5.37	7
	+		AA620437	0.16	96.0	6.02	9.62	3.03	5.40	7

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

5.20 2	7.25 2	7.87 2	7.79 2	6.52 2	5.38 2	6.09 2	8.76 2	7.53 2	2.86 2	8.22 2	8.94 2	7.60 2	6.64 2	9.69 2	8.71 2	7.58 2	6.22 2	5.62 2	5.26	8.09	6.29 2	5.11 2	1.55 2	8.40 2	7.45 2	5.77 2	10.01	5.36 2	4.86 2	1.55 2	9.54 2	1.87 2	7.73 2	7.68
9.92	9.75	7.62	8.57	9.04	11.46	0:30	7.34	3.15	8.19	8.58	6.81	5.43	2.32	9.1	5.22	5.30	72.7	9.55	10.89	5.86	6.03	10.22	10.45	7.72	7.30	9.02	6.27	9.62	99.5	9.78	6.51	9.35	8.12	5.70
2.89	0.97	2.44	1.57	2.30	0.97	11.41	1.63	7.03	9.66	<b>7</b> 8.0	1.87	4.55	8.54	6.78	3.50	4.53	3.92	2.24	1.21	3.39	4.97	<b>8</b> .	5.27	1.10	2.46	2.42	0.89	2.14	92.9	5.72	8.	5.83	1.17	3.60
9.00	5.99	5.98	6.98	5.96	5.93	5.93	5.91	5.90	5.90	5.88	5.88	5.86	5.83	5.82	5.81	5.80	5.80	5.80	5.79	5.78	5.76	5.76	5.76	5.74	5.74	5.73	5.72	5.71	5.69	5.68	5.68	5.68	2.68	5.66
77.7	68.8	8. %	2.99	0.55	69.0	1.85	15.84	3.18	0.36	0.76	2.56	8.00	4.03	90.0	43.39	10.85	32.21	4.83	0.22	13.63	0.16	0.59	4.73	2.62	3.37	6.12	1.35	0.15	4.73	1.58	5.71	2.92	2.35	A 0.8
1 29	1 48	39	05.0	800	0.12	0.31	2.68	, 1,	90:0	0.13	0.43	1.37	0.69	0.01	7.47	1.87	6.55	0.83	9.0	2.36	0.03	0.10	0.82	0.46	0.59	1.07	0.24	0.03	0.83	0.28	00.1	0.51	0.41	8
A A 458546	AA465353	AA504202	N39233	AA879423	AA101173	N62868	N36008	N29918	N59866	AA778603	N73252	W94009	AA709271	H02158	AA454585	N59136	AA625662	AA465090	AA609914	T57359	H01332	AA708605	N62128	R13434	R50752	H10788	H37817	R40481	AA150777	R45952	N72274	N51498	R89363	04330430
813410	24,080	825284	27647.2	1505686	563860	289645	272888	271076	284783	1048949	246549	357531	1343468	150623	809535	287637	745360	814995	1031113	74738	147385	506483	287581	28475	38510	46862	191546	27817	504826	34063	291374	282104	196037	803783
687	3 3	\$ 4 2	8 8	487	488	489	490	167	482	493	4 49	495	96	497	498	499	200	501	205	503	\$	505	200	507	508	209	510	511	512	513	514	515	516	547

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

2	2	7	7	7	7	7	<b>7</b>	7	7	~	2	7	7	2	7	7	2	7	7	7	7	7	7	7	7	7	7	7	7	2	2	7	2	2	
6.19	2.00	8.55	11.39	2.59	5.48	6.45	7.48	5.99	5.46	2.47	7.67	5.02	8.52	6.45	8.81	6.56	7.85	1.70	7.61	6.30	7.49	7.59	44.8	5.51	8.69	4.4	7.71	8.77	5.87	7.27	6.98	8.24	5.55	6.26	
2.36	1.37	7.77	0.44	8.42	4.48	4.88	1.26	10.00	9.88	7.41	2.85	6.93	7.04	8.01	60.0	6.29	7.74	7.64	8.14	8.29	6.97	5.09	6.79	8.15	5.96	5.49	6.69	5.32	5.58	3.79	5.22	6.68	7.08	6.07	
8.40	10.55	0.55	20.5	5.85	6.88	5.49	8.06	0.80	1.43	68.9	6.12	4.68	1.02	2.10	7.57	3.62	0.87	7.11	0.67	1.75	1.86	3.64	1.09	2.62	1.61	6.63	1.84	2.06	4.66	5.05	3.91	1.13	3.39	3.66	
5.65	5.64	5.62	29:5	5.62	5.62	5.61	5.60	5.60	5.59	5.59	5.55	5.54	5.53	5.52	5.49	5.49	5.49	5.48	5.47	5.45	5.44	5.44	5.44	5.43	5.42	5.42	5.41	5.38	5.37	5.37	5.37	5.35	5.34	5.33	16
18.53	6.68	0.25	0.10	90.0	25.20	5.38	17.75	0.0	0.17	4.03	3.65	0.40	132.57	0.41	66.94	5.08	11.11	8.93	5.38	0.26	0.05	0.31	1.05	A 45		ça o	0.59	7.13	1.59	11.60	10.19	11.61	0.55	0.94	
3.28	1.19	0.05	0.05	0.01	4.49	96.0	3.17	0.01	0.03	0.72	99.0	0.07	23.98	0.0	12.19	0.93	2.02	1.63	0.98	0.05	0.01	90.0	0.19	0.08	6.18	<b>5</b> .	0.11	1.32	0.30	2.16	1.90	2.17	0.10	0.18	
AA699560	AA669218	AA757588	AA427400	W93500	R80790	H93424	AA608730	W96197	H47315	T98201	AA035580	H23983	AA670408	AA907714	AA487608	AA115559	R22308	H80685	R92812	AA428514	H96356	AA017213	R89287	AA884428	AA706829	AA598803	AA883100	AA458827	AA779383	AA450205	AA682545	AA402965	AA676840	W93682	
433474	856592	395609	770979	357344	146882	220700	950926	361688	193139	121798	471707	51465	878788	1505898	839081	491615	130845	241432	197206	773170	256720	361069	195786	1466998	451805	898097	1467751	814350	459947	789204	431245	741790	460114	357285	
518	519	520	521	522	523	524	525	526	527	528	529	530	531	532	533	534	535	536	537	538	539	540	175	542	543	<del>2</del>	545	\$ <del>7</del>	547	£	649	920	551	552	

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line; Genes upregulated 5 fold or more in at least 2 cancer cell lines.

2	74	7	7	7	2	2	7	7	8	2	7	и	7	7	7	7	C4	2	7	7	7	7	7	2	7	2	7	2	7	2	7	7	2	7	
6.39	6.03	7.16	17.7	3.90	60.6	5.20	7.18	5.27	6.95	7.50	7.02	8.43	6.43	7.68	2	9 80	5.15	6.97	7.33	<b>3</b> 5.	5.93	8.34	5.28	5.63	5.12	7.63	0.32	5.48	2.67	7.92	5.33	5.26	5.79	1.87	
5.23	1.0.1	7.63	7.20	5.91	5.86	9.61	6.76	8.00	6.68	2.95	2.97	5.07	7.32	5.49	5.99	6.91	9.14	5.27	1.00	0.80	5.12	5.30	8.40	5.03	9.02	5.29	8.82	8.12	8.65	5.51	99.9	3.76	8.28	6.10	
4.34	8.91	1.15	1.02	60.9	9.94	1.07	1.93	2.59	2.22	5.35	5.78	2.21	1.93	2.51	2.95	1.72	1.23	3.25	7.12	6.70	4.34	1.68	1.62	4.62	1.14	2.35	6.08	<b>2</b> .	0.82	1.63	3.05	6.00	0.91	7.00	
5.32	5.32	5.31	5.31	5.30	5.30	5.29	5.29	5.29	5.28	5.26	5.26	5.24	5.23	5.23	5.19	5.18	5.17	5.16	5.15	5.15	5.13	5.11	5.10	5.09	5.09	5.09	5.07	5.05	5.04	5.02	5 02	5.01	4.99	4.99	17
35.80	0.05	6.28	2.13	13.64	21.66	4.03	0.23	20.12	0.41	3.68	0.14	0.58	44.95	15.25	0.35	9.47	0.51	2.47	0.05	90.0	31.82	0.19	0.43	0.77	32.60	1.98	16.55	40.76	0.22	3.45	23.47	4.55	<b>4</b> .5	8.53	
6.73	0.01	1.18	0.40	2.57	4.09	92.0	9.0	3.81	90:0	0.70	0.03	0.11	8.80	2.92	0.07	1.83	0.10	0.48	0.01	0.01	6.20	9.	0.08	0.15	6.40	0.39	3.28	8.07	90.0	69.0	<b>4</b> .68	0.91	0.91	1.71	
AA620556	R73525	R16676	AA598668	H98822	AA455652	N21334	N57858	177840	AA628190	H59305	AA879474	AA620614	AA133166	AA169645	AA446748	AA644334	AA426561	AA136060	AA398218	W84774	N66132	AA683338	QT7777AA	AA885851	AA148505	R71689	AA962541	AA421230	AA490605	N49898	AA504272	AA129552	H61552	AA149637	
951125	156473	129664	898258	261714	813997	265102	247089	108395	1055831	204299	1505735	1048685	490753	594517	784126	845602	763146	490060	726658	415806	278483	1293016	448920	1500388	491524	155575	1602209	739247	824109	243653	825411	564803	206052	504279	
553	554	555	556	557	558	558	260	261	925	563	564	565	999	567	999	569	920	571	572	573	574	575	929	211	278	579	280	581	285	583	584	585	586	287	

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

7	7	7	7	61	74	2	2	8	2	7	8	7	7	2	2	2	7	7	7	7	7	7	63	2	7	7	7	7	7	8	64	7	7	7	
6.61	4.03	5.79	0.97	5.65	5.61	6.75	6.79	2.27	6.33	6.25	5.46	7.16	<b>2</b> .90	6.59	1.00	5.64	5.32	1.00	6.60	7.20	5.62	5.93	6.25	6.47	6.37	8.48	6.21	6.12	5.01	5.80	<del>1</del> .	3.64	5.38	6.39	
6.33	5.47	8.07	5.74	8.14	5.83	6.08	6.72	7.27	7.99	6.45	7.62	5.39	5.34	5.47	5.04	5.49	8.39	5.50	5.68	5.26	6.24	6.79	6.82	6.47	5.38	90.0	5.74	5.51	3.86	1.33	5.90	5.23	2.63	5.33	
1.99	5.36	0.98	8.10	0.97	3.32	1.91	1.20	5.15	0.37	1.97	1.57	2.09	3.38	2.45	8.44	3.32	0.71	7.91	2.12	1.94	2.51	1.60	1.18	1.30	2.49	5.66	2.27	2.52	5.15	6.88	6.62	5.09	5.94	2.19	
4.97	4.96	4.95	46.4	4.92	4.92	4.91	4.90	4.90	4.90	4.89	4.88	4.80	4.87	4.84	4.83	19.4	4.80	4.80	4.80	4.80	4.79	4.77	4.75	4.75	4.75	4.74	4.74	4.72	4.67	4.67	4.65	4.65	4.65	4.63	18
1.93	0.75	6.34	0.05	8.75	11.53	0:30	2.11	0.08	8.32	0.71	1.49	34.71	2.66	10.43	0.05	0.90	13.99	90.0	92.90	7.55	24.62	3.4	1.68	09:0	4.36	16.44	0.31	4.54	3.48	0.12	0.27	1.69	1.82	4.18	
0.39	0.15	1.28	0.0	1.78	2.34	90:0	0.43	20.0	1.70	0.14	0.30	7.11	0.54	2.16	0.01	0.19	2.91	0.01	19.35	1.57	5.14	0.72	0.35	0.13	0.92	3.47	90.0	96.0	0.74	0.03	90:0	0.36	0.39	0.90	
NZOABO	AA626784	R44346	H39560	AA478542	AA443969	R82041	N56982	AA114106	H16743	AA904969	R55750	AA035384	AA418918	N62372	AA454554	AA156599	AA064627	R43271	AA456931	N31641	AA131526	R54846	H55915	AA521411	R00275	AA779165	AA700415	H05768	T66936	AA701587	AA676590	AA046321	H08796	AA443094	
264162	877767	34294	186132	784772	757206	147925	277545	563201	50114	1504152	40721	471598	767994	290560	809508	502296	382451	32517	838568	258966	503715	154472	204129	826089	123255	453005	460553	43826	66406	432732	882488	376764	46097	809466	
888	585	280	59.	592	593	594	585	236	269	598	288	600	601	602	603	604	605	909	209	809	609	610	611	612	613	614	615	616	617	618	619	620	621	622	

Table 2-11; Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

8	8	2	7	~	7	7	7	7	7	7	7	7	7	84	7	2	7	8	7	2	7	7	7	8	2	7	7	7	7	2	7	8	7	7	
5.88	5.45	6.25	5.98	5.05	7.00	2.28	5.85	5.23	5.23	2.00	5.54	5.34	5.12	5.56	6.95	5.02	5.57	3.12	6.78	1.71	5.93	5.98	5.87	6.76	5.17	5.29	5.41	6.30	5.23	5.54	5.26	5.86	0.44	6.47	
5.63	5.08	5.20	5.24	5.72	0.75	5.06	6.62	7.47	7.39	5.53	5.84	5.08	6.30	2.55	5.28	6.17	96.9	5.13	5.09	5.25	6.24	5.02	5.71	5.63	2.84	0.22	5.60	0.31	5.46	5.14	6.04	5.25	6.96	5.51	
2.36	3.28	2.32	2.65	2.99	5.99	6.34	1.15	0.93	0.99	1.08	2.21	3.16	2.12	5.38	1.27	2.26	0.0	5.20	1.56	6.45	1.21	2.31	1.72	0.87	5.24	7.74	2.19	6.57	2.44	2.38	1.74	1.92	5.57	0.99	
4.62	4.60	4.59	4.59	4.59	4.58	4.56	4.54	4.5 4.5	4.54	45.5	4.53	4.53	4.52	4.50	4.50	4.49	4.48	4.48	4.48	4.47	4.46	4.4	4.44	4.42	4.42	4.42	4.40	4.39	4.38	4.35	4.35	4.34	4.33	4.32	19
5.17	11.07	15.49	4.45	4.50	90.0	12.03	13.59	5.05	2.67	0.38	2.96	13.59	40.40	2.71	6.39	4.53	1.60	10.51	0.32	1.09	0.53	90.0	0.49	1.88	90.0	7.57	57.53	0.14	0.84	0.73	0.40	0.42	0.10	1.07	
1 12	2.40	3.37	0.97	0.98	0.01	2.64	2.98	1.1	69.0	0.08	9.65	300	8.95	09:0	1.42	1.01	0.36	2.35	0.07	0.24	0.12	0.01	0.11	0.42	0.02	1.71	13.07	9.03	0.19	0.17	0.09	0.10	0.02	0.25	
A 0 6 7 5 8 4 7	AA173926	AA598797	AA460521	AA284291	AA629251	AA676296	H98655	AA488979	N51367	AA704858	R24451	AA157787	AA421270	AA885096	H58542	T58434	N29624	AA454947	AA778562	H22854	AA883660	H20743	AA904776	R89765	R06372	AA398355	T90621	AA404338	AA112979	AA884382	AA626247	AA625791	AA443300	AA205320	
745102	595109	898083	796606	327245	744044	882355	261567	824728	283208	453229	132954	588840	731031	1468466	206272	75650	257312	814765	1048899	51700	1466599	51328	1504005	194811	126239	726858	110772	758302	530545	1461715	745523	744911	784589	647816	
673	524	529	628	627	628	628	630	631	523	633	834	635	636	637	638	639	64	2	275	643	3	645	8	<b>64</b>	878	200	650	651	652	653	<b>3</b> 5	655	929	159	

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

7	7	7	7	7	7	7	7	7	7	7	7	8	7	2	7	7	7	7	7	7	7
5.62	5.79	6.07	5.36	5.76	5.87	5.58	5.36	5.41	5.49	5.65	<b>6</b> .01	5.93	5.55	1.12	5.01	0.56	5.58	5.46	0.55	5.01	5.63
5.78	5.77	5.61	5.44	5.55	1.07	5.50	5.73	1.63	6.05	5.34	5.73	5.12	5.71	5.23	1.78	5.82	5.04	5.12	5.69	5.60	5.56
1.45	1.28	1.08	1.91	1.25	5.58	1.40	1.38	5.38	0.84	1.58	0.57	1.23	76.0	5.83	5.25	5.37	1.13	1.12	5.46	1.01	0.43
4.28	4.28	4.25	4.24	4.19	4.17	4.16	4.16	4.14	4.13	4.12	4.10	4.09	4.08	4.08	4.01	3.92	3.91	3.90	3.90	3.87	3.87
0.39	0.27	6.07	11.37	8.82	5.73	11.43	0.36	1.69	9.76	0.57	9.78	0.83	4.55	4.25	7.30	0.07	10.81	2.97	60:0	1.56	7.56
60.0	90'0	1,43	2.68	2.10	1.37	2.75	0.09	0.41	2.36	0.14	2.38	0.20	1.12	9.	1.82	0.02	2.77	0.76	0.02	0.40	1.95
AA437245	H08227	NSOROR	AA465692	N35579	B60949	N50935	AA883755	AA708431	AA487206	R07012	AA633993	AA862814	T54527	T62552	N79230	H79566	AA676598	H84915	N58136	H47089	AA045340
757463	45514	280909	814906	272238	42400	281103	1466824	505997	841287	128695	858292	1469211	73659	79726	292388	239661	882483	249517	247587	193420	487141
659	960	98	862	993	799	665	989	667	999	609	670	67.1	672	673	674	675	9/9	21.2	678	679	680
	757463 AA437246 0.09 0.39 4.28 1.45 5.78	757463 AA437245 0.09 0.39 4.28 1.45 5.78 45514 H08227 0.06 0.27 4.28 1.28 5.77	757463 AA437245 0.09 0.39 4.28 1.45 5.78 45514 H08227 0.06 0.27 4.28 1.28 5.77 280909 NAGRAPA 1.43 6.07 4.25 1.08 5.61	757463 AA437245 0.09 0.39 4.28 1.45 5.78 45514 H08227 0.06 0.27 4.28 1.28 5.77 280909 N50828 1.43 6.07 4.25 1.08 5.61 814906 AA46569 2.68 11.37 4.24 1.91 5.44	757463         AA437245         0.09         0.39         4.28         1.45         5.78           45514         H08227         0.06         0.27         4.28         1.28         5.77           280909         N50828         1.43         6.07         4.25         1.08         5.61           814906         AA465692         2.68         11.37         4.24         1.91         5.44           272238         N35579         2.10         8.82         4.19         1.25         5.55	757463 AA437245 0.09 0.39 4.28 1.45 5.78 45514 H08227 0.06 0.27 4.28 1.28 5.77 280309 N50828 1.43 6.07 4.25 1.08 5.61 814906 AA465692 2.68 11.37 4.24 1.91 5.44 272238 N35579 2.10 8.82 4.19 1.25 5.55 4240 BA0349 1.37 5.73 4.17 5.58 1.07	757463         AA437245         0.09         0.39         4.28         1.45         5.78           45514         H08227         0.06         0.27         4.28         1.28         5.77           280909         N50826         1.43         6.07         4.25         1.08         5.61           814906         AA465692         2.68         11.37         4.24         1.91         5.44           272238         N35579         2.10         8.82         4.19         1.25         5.55           42400         R60949         1.37         5.73         4.17         5.58         1.07           281103         N5035         2.75         11.43         4.16         1.40         5.50	757463         AA437246         0.09         0.39         4.28         1.45         5.78           45514         H08227         0.06         0.27         4.28         1.28         5.77           280909         N50828         1.43         6.07         4.25         1.08         5.51           814906         AA465692         2.68         11.37         4.24         1.91         5.44           27238         N35579         2.10         8.82         4.19         1.25         5.55           42400         R60949         1.37         5.73         4.17         5.58         1.07           281103         N50935         2.75         11.43         4.16         1.40         5.50           1466824         AA883765         0.09         0.36         4.16         1.38         5.73	757463         AA437245         0.09         0.39         4.28         1.45         5.78           45514         H08227         0.06         0.27         4.28         1.28         5.77           280909         N50828         1.43         6.07         4.25         1.08         5.51           814906         AA465692         2.68         11.37         4.24         1.91         5.44           27238         N3579         2.10         8.82         4.19         1.25         5.55           42400         R60949         1.37         5.73         4.17         5.58         1.07           281103         N50935         2.75         11.43         4.16         1.40         5.50           1466824         AA83765         0.09         0.36         4.16         1.38         5.73           505997         AA708431         0.41         1.69         4.14         5.38         1.63	757463         AA437245         0.09         0.39         4.28         1.45         5.78           45514         H08227         0.06         0.27         4.28         1.28         5.77           280909         N50828         1.43         6.07         4.25         1.08         5.61           814906         AA465692         2.68         11.37         4.24         1.91         5.44           272238         N35579         2.10         8.82         4.19         1.25         5.55           42400         R60949         1.37         5.73         4.17         5.58         1.07           281103         N50935         2.75         11.43         4.16         1.40         5.50           1466824         AA883766         0.09         0.36         4.14         5.38         1.63           50597         AA708431         0.41         1.69         4.14         5.38         1.63           841287         AA887706         2.36         9.76         4.13         0.84         6.05	757463         AA437245         0.09         0.39         4.28         1.45         5.78           45514         H08227         0.06         0.27         4.28         1.28         5.77           280909         N50228         1.43         6.07         4.25         1.08         5.61           281906         AA465692         2.68         11.37         4.24         1.91         5.44           272238         N35579         2.10         8.82         4.19         1.25         5.55           42400         R60949         1.37         5.73         4.17         5.58         1.07           281103         N50935         2.75         11.43         4.16         1.40         5.50           1466824         AA883766         0.09         0.36         4.16         1.38         5.73           605997         AA708431         0.41         1.69         4.14         5.38         1.63           841287         AA487206         2.36         9.76         4.13         0.84         6.05           126655         R07012         0.14         0.67         4.12         1.50         5.14	757463         AA437245         0.09         0.39         4.28         1.45         5.78           45514         H08227         0.06         0.27         4.28         1.28         5.77           280909         N50826         1.43         6.07         4.25         1.08         5.61           272236         AA465692         2.68         11.37         4.24         1.91         5.44           272236         AA65692         2.10         8.82         4.19         1.25         5.55           42400         R60949         1.37         5.73         4.17         5.58         1.07           281103         N50355         2.75         11.43         4.16         1.40         5.50           1466824         AA833765         0.09         0.36         4.16         1.30         5.73           505997         AA708431         0.41         1.69         4.14         5.38         1.63           841287         AA487206         2.36         9.76         4.12         1.59         5.14           858292         AA633993         2.38         9.78         4.10         0.57         5.73	757463         AA437246         0.09         0.39         4.28         1.45         5.78           45514         H08227         0.06         0.27         4.28         1.28         5.77           280909         N50828         1.43         6.07         4.25         1.08         5.51           814906         AA465692         2.68         11.37         4.24         1.91         5.44           27238         N3579         2.10         8.82         4.19         1.25         5.55           42400         R60949         1.37         5.73         4.17         5.58         1.07           281103         N50335         2.75         11.43         4.16         1.40         5.50           1466824         AA83756         0.09         0.36         4.16         1.38         5.73           1466824         AA483756         0.09         0.36         4.16         1.38         5.73           81287         AA487206         2.36         9.76         4.13         0.84         6.05           126655         RA633993         2.38         9.76         4.13         0.57         5.73           1469211         AA862814         0.20	757463         AA437246         0.09         0.39         4.28         1.45         5.78           45514         H08227         0.06         0.27         4.28         1.28         5.77           280909         N50828         1.43         6.07         4.25         1.08         5.51           814906         AA465692         2.68         11.37         4.24         1.91         5.44           272238         N35579         2.10         8.82         4.19         1.25         5.55           42400         R60949         1.37         5.73         4.17         5.58         1.07           281103         N50935         2.75         11.43         4.16         1.40         5.55           1466824         AA883765         0.09         0.36         4.16         1.38         5.73           1466824         AA487206         2.36         9.76         4.14         5.38         1.63           505997         AA487206         2.36         9.76         4.14         5.38         1.63           126655         AA633993         2.38         9.76         4.12         1.56         5.74           8882929         AA633993         2.38	757463         AA437245         0.09         0.39         4.28         1.45         5.78           45514         H08227         0.06         0.27         4.28         1.28         5.77           280909         N50828         1.43         6.07         4.25         1.08         5.61           281906         AA465692         2.68         11.37         4.24         1.91         5.44           272238         N35579         2.10         8.82         4.19         1.25         5.55           42400         R60949         1.37         5.73         4.17         5.58         1.07           281103         N50935         2.75         11.43         4.16         1.40         5.55           1466824         AA883766         0.09         0.36         4.16         1.38         5.73           1466824         AA883766         0.09         0.36         4.16         1.38         5.73           841287         AA487206         2.36         9.76         4.14         5.38         1.63           126695         R07012         0.14         0.57         4.12         1.58         5.14           858292         AA633993         2.38	757463         AA437245         0.09         0.39         4.28         1.45         5.78           45514         H08227         0.06         0.27         4.28         1.28         5.77           280309         N50228         1.43         6.07         4.25         1.08         5.51           219050         AA65692         2.68         11.37         4.24         1.91         5.44           272238         N35579         2.10         8.82         4.19         1.25         5.55           42400         R60349         1.37         5.73         4.17         5.58         1.07           281103         N50355         2.75         11.43         4.16         1.40         5.56           1466824         AA883766         0.09         0.36         4.16         1.40         5.50           1466824         AA883766         0.09         0.36         4.16         1.38         5.73           841287         AA708431         0.41         1.69         4.14         5.38         1.63           126695         R07012         0.14         0.57         4.12         1.58         5.14           856292         AA633993         2.38	757463         AA437245         0.09         0.39         4.28         1.45         5.78           45514         H08227         0.06         0.27         4.28         1.28         5.77           280309         N50828         1.43         6.07         4.25         1.08         5.61           281906         AA465692         2.68         11.37         4.24         1.91         5.44           272236         N35579         2.10         8.82         4.19         1.25         5.55           42400         R60949         1.37         5.73         4.17         5.58         1.07           281103         N50355         2.75         11.43         4.16         1.40         5.56           1466824         AA843766         0.09         0.36         4.16         1.30         5.73           1466824         AA843766         0.09         0.36         4.16         1.38         5.73           841287         AA708431         0.41         1.69         4.14         5.38         1.63           126955         R07012         0.14         0.57         4.12         1.59         5.73           1469211         AA862814         0.20	757463         AAA37246         0.09         0.39         4.28         1.45         5.78           45514         H08227         0.06         0.27         4.28         1.28         5.77           280309         N50228         1.43         6.07         4.25         1.08         5.51           280309         N50228         1.43         6.07         4.25         1.08         5.51           281030         N50328         2.10         8.82         4.19         1.25         5.55           272238         N35579         2.10         8.82         4.19         1.25         5.55           272238         N35035         2.75         11.43         4.17         5.58         1.07           281103         N50935         2.75         11.43         4.16         1.40         5.56           1468824         AA883766         0.09         0.36         4.16         1.38         5.73           44128         AA88284         0.41         1.69         4.14         5.38         1.63           44128         AA88284         0.20         0.83         4.09         0.57         5.73           146927         1.12         4.55         4.	75763         AA437245         0.09         0.39         4.28         145         5.78           45514         H08227         0.06         0.27         4.28         1.28         5.77           280909         N50828         1.43         6.07         4.25         1.08         5.51           280909         N50828         1.43         6.07         4.25         1.08         5.77           2814906         AA465692         2.68         11.37         4.24         1.91         5.44           272236         N35579         2.10         8.82         4.19         1.25         5.55           22400         R60949         1.37         5.73         4.17         5.58         1.07           281103         N50935         2.75         11.43         4.16         1.40         5.56           1468824         AA883766         0.09         0.36         4.16         1.30         5.73           841287         AA481206         2.36         9.76         4.14         5.38         1.63           848292         AA633993         2.38         9.76         4.13         0.84         6.05           146921         AA632993         2.38	757463         AA437246         0.09         0.39         4.28         1.45         5.78           45514         H08227         0.06         0.27         4.28         1.28         5.77           280909         N50828         1.43         6.07         4.25         1.08         5.51           814906         AA465692         2.68         11.37         4.24         1.91         5.44           27236         N3579         2.10         8.82         4.19         1.25         5.55           27208         N3579         1.37         5.73         4.17         5.58         1.07           28400         R60349         1.37         5.73         4.16         1.40         5.56           146824         AA483766         0.09         0.36         4.16         1.39         5.73           446824         AA487206         2.36         9.76         4.13         0.84         6.05           60597         AA487206         2.36         9.76         4.13         0.84         6.05           841287         AA487206         0.14         0.57         4.12         1.58         5.73           126692         AA65814         0.14 <th< th=""><th>650         75740         AA437246         0.09         0.39         4.28         1445         5.78         5.62         2           660         45514         H082724         0.06         0.27         4.28         128         5.77         5.79         2           661         25614         H08228         1.43         6.07         4.25         1.08         5.71         6.79         2           662         814906         AA465622         2.68         11.37         4.24         1.91         5.44         5.36         2           663         27220         N35579         2.10         8.82         4.19         1.25         5.55         5.76         2           664         42400         R60949         1.37         4.17         5.58         1.07         5.87         5.76         2           665         281103         N5035         2.75         11.43         4.16         1.39         5.59         5.73         5.76         5.73         5.73         5.74         5.76         2           665         281103         N50350         2.75         11.43         4.16         1.39         5.73         5.73         5.73         5.74</th></th<>	650         75740         AA437246         0.09         0.39         4.28         1445         5.78         5.62         2           660         45514         H082724         0.06         0.27         4.28         128         5.77         5.79         2           661         25614         H08228         1.43         6.07         4.25         1.08         5.71         6.79         2           662         814906         AA465622         2.68         11.37         4.24         1.91         5.44         5.36         2           663         27220         N35579         2.10         8.82         4.19         1.25         5.55         5.76         2           664         42400         R60949         1.37         4.17         5.58         1.07         5.87         5.76         2           665         281103         N5035         2.75         11.43         4.16         1.39         5.59         5.73         5.76         5.73         5.73         5.74         5.76         2           665         281103         N50350         2.75         11.43         4.16         1.39         5.73         5.73         5.73         5.74

Table 2-12: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line. Genes upregulated 10 fold or more in at least 2 cancer cell lines.

Table 2-12: Prostate Cancer Cell Lines vs. Normal

Number         Values           AA725564         0.01         1.12           AA47079         0.01         0.66           AA460463         0.01         0.35           AA406036         0.01         0.50           AA406036         0.01         0.50           AA406036         0.01         0.50           AA406036         0.01         0.32           AA406036         0.01         0.32           AA456035         0.01         0.32           H09818         0.01         0.27           H09818         0.05         17.34           NNZ115         0.23         6.85           H02294         0.48         12.13           H02294         0.48         12.53           R59556         0.01         0.17           AA02840         0.40         10.00           AA778675         0.38         9.30           AA629838         0.01         0.24           AAA28674         0.17         2.95           AA538374         1.21         16.52           AA448350         0.01         9.14           AA44846         0.01         1.70           AA44646<	Order	MAGE	Gen Bank Accession	MPM1(PrEC)	Ave Cancer Cell Lines	Fold (Ave Cell	Fold(MPM2/M PM1)	Fold(MPM2/M Fold(MPM271/MP Fold(MPM272/M # of Cancer PM1) M1) Cell Lines	Fold(MPM272/M PM1)	# of Cancer Cell Lines
144772         AA725864         UOI         1.12         1.2.2         1.2.2         1.4.4772         1.4.4772         1.4.4772         1.4.4772         1.4.4772         1.4.4773         1.4.4773         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774         1.4.4774 <th>,</th> <th></th> <th>Number</th> <th>č</th> <th>Values</th> <th>112.48</th> <th>13.45</th> <th>301.13</th> <th>22.85</th> <th>n</th>	,		Number	č	Values	112.48	13.45	301.13	22.85	n
784288         AAA47079         0.01         0.05         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.03         0.04         0.03         0.04         0.03         0.04         0.03         0.04         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03         1.03	-	1343/32	AA725564		71.1	06.30	45.84	106.60	43.45	က
796559         AA460463         0.01         0.35         0.557         0.10         0.20           7305655         AA460466         0.01         0.50         55.37         0.10         0.10           743016         AA460636         0.01         0.50         50.47         15.70         17.32           520307         AA460636         0.01         0.50         36.21         45.36         56.36         20.37           626307         AA460636         0.07         3.16         45.16         91.62         20.97           81270         AA466036         0.01         0.30         37.18         16.25         80.79           81270         AA466036         0.01         0.32         35.79         12.25         87.45           9028         H1786         0.01         0.32         35.79         12.25         80.79           291057         M2715         0.22         0.59         22.44         39.44           4876         H0381         0.02         0.59         22.71         13.82         30.16           4889         H0381         0.02         0.59         22.71         13.82         30.16           41516         H03224         <	~	784296	AA447079	0.04	0.56	00° 50	£	138.91	35.70	က
1558655         AA978561         0.08         4.36         54.32         97.06         77.32           743016         AA406036         0.01         0.50         50.47         15.70         123.80           33076         AA406036         0.01         0.50         50.47         15.70         123.80           626300         AA189106         0.07         3.16         45.16         91.62         20.97           626300         AA189106         0.07         3.16         45.16         91.62         20.97           626300         AA18910         0.07         0.30         37.18         16.25         80.79           487861         AA04436         0.01         0.32         35.79         12.25         87.45           281057         AA04436         0.02         0.59         32.71         12.25         87.45           281057         AA04436         0.01         0.22         25.94         38.94         38.44           281057         AA04436         0.01         0.27         27.21         13.86         30.16           45896         H03818         0.05         0.17         25.59         33.88         30.16           41945         R585	3	796569	AA460453	0.01	0.35	63.57	16.10	16.901	47.07	
743016         AAAG6036         0.01         0.50         50.47         15.70         123.80           33076         RA406036         0.01         36.21         45.35         56.36         29.19           626390         AA489106         0.07         3.16         45.16         91.62         20.97           812170         AA456035         0.01         0.30         37.18         16.25         80.73           812170         AA456035         0.01         0.32         35.79         12.25         80.74           48761         AA456036         0.01         0.32         35.79         12.25         87.45           291057         ANZ115         0.23         23.79         12.25         87.45           291057         ANZ115         0.23         23.73         24.48         38.30           203014         H93818         0.01         0.27         27.21         13.82         36.04           46886         H03818         0.05         12.1         25.99         33.86         13.60           41595         R59556         0.01         0.17         25.59         3.86         13.61           41596         AA03840         0.40         10.00 <td>4</td> <td>1558655</td> <td>AA976561</td> <td>90.0</td> <td>4.36</td> <td>54 32</td> <td>97.06</td> <td>17.92</td> <td>18.74</td> <td>n (</td>	4	1558655	AA976561	90.0	4.36	54 32	97.06	17.92	18.74	n (
33076         R44048         0.80         38.21         45.35         56.36         29.19           626390         AA189106         0.07         3.16         45.16         91.62         20.97           812170         AA456035         0.01         0.30         37.18         16.25         80.79           80288         H17886         0.01         0.32         35.79         12.25         87.45           487861         AAA45635         0.02         0.59         32.57         29.44         39.44           291057         N72115         0.23         0.59         22.919         24.48         38.30           263014         H99813         0.02         0.59         22.919         24.48         38.30           263014         H99813         0.01         0.27         27.21         13.07         21.59           46896         H09818         0.05         1.21         25.96         33.86         30.16           45896         H09818         0.05         1.21         25.96         33.86         30.16           41595         R5956         0.01         0.07         1.25         27.21         13.86         30.16           415961	. M	743016	AA406036	0.01	0.50	50.47	15.70	123.80	11.93	m ·
626339         AA489106         0.07         3.16         45.16         91.62         20.97           812179         AA426035         0.01         0.30         37.18         16.25         80.79           812170         AA436035         0.01         0.32         35.79         12.25         87.45           50286         H17866         0.01         0.32         35.79         12.25         87.45           487861         AA426436         0.02         0.59         32.57         29.44         39.44           291057         N72115         0.22         0.59         32.91         24.48         38.30           263014         H99813         0.02         0.27         27.21         13.82         36.04           46896         H99818         0.01         0.27         27.21         13.82         30.16           46896         H99818         0.05         1.21         25.96         33.89         13.26           45896         H90818         0.05         0.17         25.59         33.89         13.26           41595         R59556         0.01         0.17         25.59         33.89         13.64           41594         AA70002		33076	R44048	0.80	36.21	45.35	56.36	29.19	50.51	ო
912170         AA456035         0.01         0.30         37.18         16.25         80.79           50288         H17886         0.01         0.32         35.79         12.25         87.45           291057         N72115         0.23         0.59         32.57         29.48         39.44           291057         N72115         0.23         6.85         29.19         24.48         38.30           263014         H99813         0.63         17.34         27.31         19.07         21.59           303180         N92764         0.01         0.27         27.21         13.82         38.04           46896         H09818         0.05         1.21         25.96         33.86         30.16           45896         H09818         0.05         1.21         25.96         33.86         30.16           45896         H09818         0.05         1.21         25.96         33.89         30.16           45896         H09818         0.05         1.21         25.96         33.89         13.26           41594         R59556         0.01         0.17         25.59         33.89         13.66           43002         AA778675         <	) h	626390	44189106	20 0	3.16	45.16	91.62	20.97	22.90	က
50288         H17886         0.01         0.32         35.79         12.25         87.45           487861         AAQ45436         0.02         0.59         32.57         29.44         39.44           291057         N72115         0.23         0.59         32.57         29.48         38.30           29057         N72115         0.23         0.53         17.34         27.31         19.07         21.59           263014         H99813         0.05         1.21         25.96         33.86         30.16           46896         H09818         0.05         1.21         25.96         33.86         30.16           45896         H09818         0.05         1.21         25.96         33.86         30.16           45896         H09818         0.05         0.17         25.59         33.86         30.16           41595         R5956         0.01         0.17         25.59         33.86         30.16           41596         AAC28483         0.01         0.07         2.480         34.18         13.26           43002         AAC28433         0.01         0.24         24.29         34.18         16.26           84879	- «	812170	AA456035	5	0:30	37.18	16.25	80.79	14.51	က
487861         AAA445436         O.23         6.85         32.57         29.44         39.44           281057         N72115         0.23         6.85         29.19         24.48         38.30           281057         N72115         0.23         6.85         29.19         24.48         38.30           283014         H93813         0.63         17.34         27.21         13.82         36.04           46896         H03816         0.05         1.21         25.96         33.86         30.16           45896         H03214         0.48         1.2.53         25.93         38.89         13.26           41595         R58556         0.01         0.17         25.59         12.08         52.20           430092         AA08840         0.40         10.00         24.80         34.18         13.26           44595         R58556         0.01         0.07         25.59         12.08         52.20           430092         AA08840         0.40         10.00         24.80         13.6         10.6           44596         AA028138         0.01         0.24         24.29         30.26         10.99           46516         AA084518	• •	80008	7888		0.32	35.79	12.25	87.45	10.68	က
24.05         AZGAZIA	, 5	487861	AA045436	200	65.0	32.57	29.44	39.44	28.82	က
203014         H99613         0.63         17.34         27.31         19.07         21.59           303180         N92764         0.01         0.27         27.21         13.82         36.04           46886         H09818         0.05         1.21         25.96         33.86         30.16           46886         H09818         0.05         1.21         25.99         33.86         30.16           41595         R59566         0.01         0.17         25.59         12.08         52.20           41595         R59566         0.01         0.17         25.59         12.08         52.20           41595         R59566         0.01         0.17         25.59         12.08         52.20           41596         AA009840         0.40         10.00         24.80         34.18         13.61           1049033         AA778675         0.38         9.30         24.51         46.38         16.26           884790         AA62948         0.01         0.24         24.29         30.26         10.09           365148         AA025142         0.16         3.52         22.24         38.20         16.36           488431         AA647441	2 =	291057	N72115	0.23	6,85	29.19	24.48	38.30	24.80	ю
303180         N92764         0.01         0.27         27.21         13.82         36.04           46896         H09818         0.05         1.21         25.96         33.86         30.16           45896         H09818         0.05         1.253         25.99         33.89         13.26           41595         R59556         0.01         0.17         25.59         12.08         52.20           430092         AA009840         0.40         10.00         24.80         34.18         13.51           1049033         AA778675         0.38         9.30         24.51         46.36         13.61           1049033         AA778675         0.38         9.30         24.51         46.36         15.26           1049033         AA778675         0.38         9.30         24.29         30.26         10.09           365149         AA025142         0.01         0.24         24.29         30.26         10.09           365149         AA025142         0.01         0.25         22.24         38.20         16.36           48431         AA047441         0.17         2.95         17.83         10.45         28.90           42627         R6095	: :	267014	F1866H	8	17.34	27.31	19.07	21.59	41.29	ന
46896         H09818         0.05         1.21         25.96         33.86         30.16           46896         H09818         0.05         1.21         25.93         38.89         13.26           41595         R59556         0.01         0.17         25.59         12.08         52.20           430092         AA009840         0.40         10.00         24.80         34.18         13.61           1049033         AA778675         0.38         9.30         24.51         46.38         16.26           864790         AA629838         0.01         0.24         24.29         34.26         10.09           365148         AA025142         0.01         0.24         24.29         30.26         10.09           365148         AA025142         0.01         0.25         22.24         38.20         16.96           51681         H22824         0.01         0.25         20.42         18.18         18.45           488431         AA047441         0.17         2.95         17.83         10.45         28.90           82869         AA48350         0.01         9.14         660.46         2.32         81.14         14.07           757165 <td>i ¢</td> <td>303180</td> <td>N92764</td> <td>0.01</td> <td>0.27</td> <td>27.21</td> <td>13.82</td> <td>36.04</td> <td>31.77</td> <td>က</td>	i ¢	303180	N92764	0.01	0.27	27.21	13.82	36.04	31.77	က
151184         H02294         0.48         12.53         25.93         38.89         13.26           41595         R59556         0.01         0.17         25.59         12.08         52.20           430092         AA009840         0.40         10.00         24.80         34.18         13.61           1049033         AA778675         0.38         9.30         24.51         46.38         16.26           884790         AA629838         0.01         0.24         24.29         30.26         10.09           385149         AA025142         0.16         3.52         22.24         38.20         16.96           51681         H22824         0.01         0.24         24.29         30.26         10.09           385149         AA025142         0.01         0.24         24.29         30.26         10.09           385149         AA025142         0.01         0.25         22.24         38.20         16.96           488431         AAA4441         0.02         0.31         18.24         10.20         24.35           42829         AA589974         1.21         16.52         13.65         13.47         11.86           42627         AA46	. 1	45896	HOORIA	\$0.0	1.21	25.96	33.86	30.16	13.87	က
41595         RS9556         0.01         0.17         25.59         12.08         52.20           430032         AA009840         0.40         10.00         24.80         34.18         13.61           1049033         AA72675         0.38         9.30         24.51         46.38         16.26           884790         AA629838         0.01         0.24         24.29         30.26         10.09           365149         AA025142         0.16         3.52         22.24         30.26         10.09           365149         AA025142         0.16         0.25         20.42         38.20         16.96           51581         H22824         0.01         0.25         20.42         18.16         18.45           1467167         AA83187         0.02         0.31         18.24         10.20         24.35           488431         AAA4441         0.17         2.95         17.83         10.45         28.90           488431         AAA4441         0.17         2.95         17.83         10.45         28.90           42627         R60995         0.02         0.32         15.94         21.48         14.07           757165         AA44646	. Æ	151184	H02294	0.48	12.53	25.93	38.89	13.26	25.64	က
430092         AA009840         0.40         10.00         24.80         34.18         13.61           1049033         AA778675         0.38         9.30         24.51         46.38         16.26           884790         AA629838         0.01         0.24         24.29         30.26         10.09           365149         AA025142         0.16         3.52         22.24         38.20         16.96           51581         H22824         0.01         0.25         20.42         18.16         18.45           1467161         AA883187         0.02         0.31         18.24         10.20         24.35           488431         AA047441         0.17         2.95         17.83         10.45         28.90           82869         T69270         0.02         0.32         15.94         21.48         14.07           82869         AA58874         1.21         16.52         13.65         13.47         11.86           42627         AA443950         0.01         9.14         660.46         2.32         61.14         1           591.07         AA46466         0.01         1.66         234.30         8.10         680.66           781036<	. <u>.</u>	41595	R59556	0.01	0.17	25.59	12.08	52.20	12.48	'n
1049033         AA778675         0.38         9.30         24.51         46.38         16.26           984790         AA629838         0.01         0.24         24.29         30.26         10.09           365148         AA025142         0.16         3.52         22.24         38.20         16.96           51681         H22824         0.01         0.25         20.42         18.16         18.45           1467161         AA883187         0.02         0.31         18.24         10.20         24.35           488431         AA047441         0.17         2.95         17.83         10.45         28.90           82865         T69270         0.02         0.32         15.94         21.48         14.07           82865         AA588974         1.21         16.52         13.65         13.47         11.86           42627         AA443950         0.01         9.14         660.46         2.32         61.14         1           757165         AA44646         0.01         1.66         234.30         8.10         680.66           781036         AA44646         0.01         1.70         228.86         10.58         674.66	2 5	430092	AA009840	0.40	10.00	24.80	34.18	13.61	28.60	က
884790         AA629838         0.01         0.24         24.29         30.26         10.09           365148         AA025142         0.16         3.52         22.24         38.20         16.96           51681         H22824         0.01         0.25         20.42         18.18         18.45           1467161         AA883187         0.02         0.31         18.24         10.20         24.35           488431         AA047441         0.17         2.95         17.83         10.45         28.90           82869         T69270         0.02         0.32         15.94         21.48         14.07           898286         AA588974         1.21         16.52         13.65         13.47         11.86           42627         R60995         0.01         9.14         660.46         2.32         61.14         1           757165         AA443950         0.01         5.71         591.67         7.33         66.37         1           781036         AA44646         0.01         1.70         228.86         10.58         674.66	. «	1049033	AA778675	0.38	9.30	24.51	46.38	16.26	10.88	3
365149         AA025142         0.16         3.52         22.24         38.20         16.96           61681         H22824         0.01         0.25         20.42         18.18         18.45           1467161         AA883187         0.02         0.31         18.24         10.20         24.35           488431         AA047441         0.17         2.95         17.83         10.45         28.90           82869         T69270         0.02         0.32         15.94         21.48         14.07           896286         AA598974         1.21         16.52         13.65         13.47         11.86           42627         R60995         0.01         9.14         660.46         2.32         61.14           757165         AA443950         0.01         5.71         591.67         7.33         66.37           781036         AA44646         0.01         1.70         228.86         10.58         674.66	2 9	884790	AA629838	0.01	0.24	24.29	30.26	10.09	32.50	က
61581         H22824         0.01         0.25         20.42         18.16         18.45           1467161         AA83187         0.02         0.31         18.24         10.20         24.35           488431         AA647441         0.17         2.95         17.83         10.45         28.90           488431         AA647441         0.17         2.95         17.83         10.45         28.90           82869         T69270         0.02         0.32         15.94         21.48         14.07           896286         AA588374         1.21         16.52         13.65         13.47         11.86           42627         R60995         0.01         9.14         660.46         2.32         61.14           757165         AA443850         0.01         5.71         591.67         7.33         66.37         1           781036         AA44646         0.01         1.70         228.86         10.58         674.66	: 5	365149	AA025142	0.18	3.52	22.24	38.20	16.96	11.55	ო
1467161         AA883187         0.02         0.31         18.24         10.20         24.35           488431         AA047441         0.17         2.95         17.83         10.45         28.90           82869         T69270         0.02         0.32         15.94         21.48         14.07           82869         AA598974         1.21         16.52         13.65         13.47         11.86           42627         R60995         0.01         9.14         660.46         2.32         61.14           757165         AA443950         0.01         5.71         591.67         7.33         66.37         1           781036         AA44646         0.01         1.70         228.86         10.58         674.66	3 5	51581	H22824	0.01	0.25	20.42	18.18	18.45	24.62	е
488431 AA047441 0.17 2.95 17.83 10.45 28.90 82869 T69270 0.02 0.32 15.94 21.48 14.07 698286 AA598974 1.21 16.52 13.65 13.47 11.86 42627 R60995 0.01 9.14 660.46 2.32 81.14 14.07 757165 AA443950 0.01 5.71 591.67 7.33 66.37 13.00 680.66 592707 AA160606 0.01 1.66 224.30 8.10 680.66 781036 AA44646 0.01 1.70 228.86 10.58 674.66	: 2	1467163	AA883187	0.02	0.31	18.24	10.20	24.35	20.16	က
82869         T69270         0.02         0.32         15.94         21.48         14.07           898286         AA598974         1.21         16.52         13.65         13.47         11.86           42627         R60995         0.01         9.14         660.46         2.32         81.14         1           757165         AA443950         0.01         5.71         591.67         7.33         66.37         1           592707         AA460608         0.01         1.66         224.30         8.10         680.66           781036         AA446446         0.01         1.70         228.86         10.58         674.66	! ह	488431	A A 0.47441	0.17	2.95	17.83	10.45	28.90	14.13	•
696286         AA588974         1.21         16.52         13.65         13.47         11.86           42627         R60995         0.01         9.14         660.46         2.32         61.14         1           757165         AA443950         0.01         5.71         591.67         7.33         66.37         1           592707         AA160608         0.01         1.66         234.30         8.10         680.66           781036         AA44646         0.01         1.70         228.86         10.58         674.66	3 7	82869	T69270	0.02	0.32	15.94	21.48	14.07	12.25	က
42627 R60995 0.01 9.14 660.46 2.32 61.14 1 1 2 1 2 2 1 1 1 1 1 1 1 1 1 1 1 1 1	. <b>.</b> .	898286	AA598974	1.21	16.52	13.65	13.47	11.86	15.62	ო
757165 AA443950 0.01 5.71 591.67 7.33 66.37 1 592707 AA160608 0.01 1.66 224.30 8.10 680.66 781036 AA44646 0.01 1.70 228.86 10.58 674.66	, e	42627	860995	0.01	9.14	660.46	2.32	61.14	1917.92	2
592707 AA160608 0.01 1.66 234.30 8.10 680.66 781036 AA44646 0.01 1.70 228.86 10.58 674.66	3 2	757165	AA443950	0.01	5.71	591.67	7.33	66.37	1701.31	7
781036 AA44646 0.01 1.70 228.86 10.58 674.66	: %	592707	AA160608	0.01	1.66	234.30	8.10	680.66	14.13	7
	g	781036	AA446446	0.01	1.70	228.86	10.58	674.66	1.35	2

Table 2-12: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line. Genes upregulated 10 fold or more in at least 2 cancer cell lines.

2	2	۲۷ ,	7	2	7	7	7	7	7	63	7	7	7	7	7	7	7	7	7	7	7	7	7	64	7	7	7	7	7	7	7	77	7	7	
10.82	146.14	105.59	12.54	13.76	99.88	11.76	70.45	123.70	0.72	12.17	5.49	91.51	6.93	11.05	19.33	12.68	51.22	9,41	11.01	55.73	48.14	43.19	14.72	11.03	11.15	52.45	36.33	63.66	14.11	10.31	16.63	11.62	31.35	45.50	
570.75	140.40	150.46	239.88	229.57	95.72	168.10	4.69	51.92	88.50	139.36	44.77	56.58	101.07	109.53	99.20	88.28	52.03	73.72	82.72	34.54	30.24	36.47	65.78	70.25	69.04	27.83	35.03	11.75	58.23	66.04	54.64	60.05	41.17	26.82	
8.10	0.80	0.89	3.09	1.42	0.87	3.73	104.24	1.03	78.84	0.63	99.52	0.91	25.03	4.59	4.81	5.91	0.93	20.19	5.81	72.0	9.10	7.73	5.99	0.74	1.21	0.69	7.36	2.28	5.18	0.59	4.46	3.61	2.32	2.27	
196.56	95.78	85.65	85.17	81.58	65.49	61.20	69.79	58.88	56.02	50.72	49.93	49.67	44.34	41.72	41.11	35.62	34.73	¥.	33.18	30.35	29.16	29.13	28.83	27.34	27.13	26.99	26.24	25.90	25.84	25.65	25.24	25.09	24.95	24.86	2
1.28	1.93	6.83	0.51	0.82	5.16	0.61	26.93	2.87	12.41	1.10	5.02	13.02	4.0	0.42	0.41	0.36	8.56	0.25	0.33	7.89	0.40	0.41	0.44	0.27	0.27	14.38	0.16	14.15	0.24	0.39	0.25	0.17	0.25	24.91	
0.01	0.02	90:0	0.01	0.0	90:0	0.01	0.45	9.03	0.22	0.02	0.10	0.26	0.0	0.01	0.01	0.01	0.25	0.01	0.01	0.26	0.01	0.01	0.02	0.01	0.01	0.53	0.01	0.55	0.01	0.02	0.0	0.01	0.01	1.00	
H95633	H56453	AA485214	AA459917	W80611	AA419486	W94289	R23270	AA251784	W73144	R44741	R51617	AA504844	R09747	AA437126	AA810225	H94482	R91566	N30098	AA603832	AA485052	AA863292	177729	AA789328	R15740	AA460274	H50655	AA418903	N47445	R49555	N52767	AA010383	AA236986	W94486	H19826	
220851	203878	815794	795627	347182	746152	346942	131452	684879	344589	33827	39442	825740	128083	757368	1367900	243181	196544	256911	1030613	815861	1455835	108378	1391682	53039	796549	194318	768056	280699	38344	283379	430255	687972	357884	172517	
30	3 5	35	33	ಸ	35	36	37	38	39	9	¥	42	<b>t</b>	1	45	46	47	84	6	90	5	. 25	53	ま	55	88	22	58	20	.09	61	62	63	2	

Table 2-12: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line. Genes upregulated 10 fold or more in at least 2 cancer cell lines.

														٠																						
·	7	7	7	8	7	7	7	7	7	2	7	8	7	7	7	7	64	2	7	8	2	7	7	7	7	74	7	7	7	7	7	7	7	7	7	
•	67.11	43.13	96.6	48.25	26.43	25.34	31.76	11.14	<b>54</b> .20	17.82	3.08	47.62	22.42	17.48	9.30	28.27	20.17	45.41	40.27	25.28	1.08	43.06	10.71	16.22	10.73	16.89	12.21	34.04	31.84	34.18	33.48	26.82	28.36	25.15	19.22	
:	60.48	28.65	17.65	21.91	34.50	42.67	30.37	51.95	10.05	1.50	23.72	10.88	33.42	42.76	30.57	30.54	38.72	10.31	11.18	30.81	14.31	10.12	38.83	34.15	33.82	8.91	37.48	11.78	10.67	1.51	5.09	15.89	16.52	21.09	22.78	
;	1.76	0.95	44.37	0.87	8.11	1.0	4.98	3.82	0.88	39.83	37.62	4.40	5.36	0.93	20.94	1.29	1.00	2.79	7.04	2.35	41.41	3.39	4.49	1.97	7.62	25.95	1.87	5.64	8.16	14.65	10.47	5.81	2.63	0.69	4.69	
	24.50	24.24	23.97	23.68	23.01	23.00	22.37	22.30	21.71	21.70	21.44	20.97	20.40	20.39	20.27	20.04	19.97	19.50	19.50	19.47	18.93	18.85	18.01	17.44	17.36	17.25	17.19	17.15	16.89	16.78	16.35	16.17	15.84	15.64	15.57	ო
	0.24	7.41	0.16	3.51	0.16	0.23	0.16	3.74	3.65	0.14	40.77	19.16	0.20	6.19	2.46	3.94	0.20	0.20	0.19	3.30	21.01	0.17	0.35	0.17	0.21	33.78	1.06	0.63	0.17	0.11	0.16	11.31	2.94	5.98	0.30	
	0.01	0.31	0.01	0.15	100	0.0	0.0	0.17	0.17	0.01	1.90	0.91	0.01	0.30	0.12	0.20	0.0	0.0	0.01	0.17	1:1	0.01	0.05	0.0	0.01	1.96	90.0	9.0	0.01	0.01	0.01	0.70	0.19	0.38	0.02	
	H96647	AA490945	R39191	AA504625	N63057	T70198	AA630328	R44936	AA491261	H10072	T61938	N32804	N32542	AA255954	H22563	R95684	AA461508	R86970	R54036	H28091	T65736	R54492	AA057185	AA001376	AA701315	T49355	AA857163	N47012	AA676998	AA481481	W92263	W93717	AA504838	AA417713	AA045175	
	251806	824025	23345	825606	284714	81050	855910	34204	824270	46843	2,000	259884	270875	686552	51916	199337	795631	197265	40010	162310	80338	39722	489076	361899	435099	67440	1410444	280155	454083	756629	358990	357373	825726	746232	487766	
	65	99	29	89	3 6	2 2	: =	. 22	: 22	. 7.	75	92	1	. 28	92	80		. F	83	3 3	. <del>8</del>	8	87	8 8	28	8	2	8	8	3	92	96	26	98	66	

Table 2-12: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line. Genes upregulated 10 fold or more in at least 2 cancer cell lines.

2	7	~	2	7	2	2	7	2	~	۰ ،		<b>v)</b> (	7 (	7 (	7	<b>73</b>	71	2	7	63	7	~	2	7	~	2	7	~	7	74	2	73	71	7	7	
15.13	18.34	15.00	22.27	15.54	23.25	28.45	27.66	4.48	20.74	77 7	5.15	14.71	18.2/	19.45	18.18	5.86	26.50	25.70	11.26	0.58	26.39	20.98	11.09	17.09	17.55	20.10	12.88	16.24	19.11	16.72	14.01	12.09	3.40	20.99	16.34	
28.43	26.10	24.61	16.46	3.02	18.93	4.19	11.04	18 07	27.0	07.6	70.44	15.99	21.58	12.04	18.08	17.85	1.54	12.28	21.72	14.55	0.39	14.19	23.87	13.80	15.47	14.95	18.70	17.13	13.18	8.33	20.54	19.31	20.61	10.18	16.85	
1.53	0.42	5.14	5.11	24.69	1 00	20 01	2 99	3 1	17.03	10.04	2.42	7.33	29.0	8.23	3.39	14.72	10.97	0.65	5.65	22.57	10.40	1.96	2.11	5.65	3.28	1.00	4.34	2.08	3.09	10.15	0.58	3.60	10.87	3.65	0.33	
15.03	14.95	14.92	1461	14.41	14.40	5. 7.	14.24	13.03	13.76	13.71	13.68	13.58	13.50	13.24	13.22	13.14	13.00	12.88	12.88	12.56	12.39	12.38	12.36	12.18	12.10	12.02	11.98	11.82	11.80	11.73	11.71	11.67	11.63	11.61	11.17	•
3.91	3.94	0.35	0.35	0.0	5 7	4 7	5.51	4.1/	0.13	15.63	1.31	0.17	3.95	8.63	0.26	0.23	90.0	0.13	0.17	0.13	44.0	0.12	3.72	14.39	0.31	1.69	16.82	23.53	3.89	0.31	0.18	11.24	0.20	27.64	45.0	
920	920	200	9 0.04	0.02	- 6		0.25	0.30	0.01	4.4	0.10	0.01	0.29	0.65	0.02	0.02	0.01	500	0.01	0.01	0.76	0.01	030	1.18	500	0. 4.	1,40	8	0.33 EE.0	0.03	0.02	0.96	200	2.38	0.03	! !
00463446	140041	70997	H8/140	K40434	AA4/6543	AA434388	AA883800	W90705	R56149	AA628867	R37738	AA128617	AA427719	W93379	AA491227	H19217	N52193	AA017379	N34042	A4453618	AA620697	AA022496	H10713	AA262211	AA778717	R92201	AA706901	AA477227	H09940	AA609002	AA161188	R44816	FC#F0W1	AA02682	AA160670	
044766	611100	Accid	8000c7	96797	182283	770860	1461609	418004	41406	1035796	26806	502003	771004	415089	824074	51103	28428B	361122	244050	70544R	1049185	364352	47.784	686172	1040079	195845	451649	739450	45461	1030798	591167	33839	347138	366971	592728	) 
Ş	3 3	<u> </u>	20 1	<b>SD</b> :	ž	<b>5</b>	8	107	108	109	110	111	112	113	4.	115	. <del>.</del>	÷		2 5	• ¢	<u> </u>	į į	<u> </u>	3 5	125	126	121	£ 5	\$ 5	0.55	5	<u>;</u> £	<u> </u>	3 5	•

Table 2-12: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line. Genes upregulated 10 fold or more in at least 2 cancer cell lines.

2	7	2	7	7	7	2	7	2	7	2	7	7	7	~ .	2	7	7	~	~	7	7	7	7	64	7	7	7	2	8	2	7	7	7	7	
13.80	16.62	10.20	12.02	20.10	12.92	7.28	15.60	15.64	1.65	17.42	16.30	14.06	15.31	10.88	7.37	10.89	5.79	11.50	15.44	15.97	15.06	14.44	5.24	4.33	11.30	13.51	13.08	14.35	14.67	1.10	16.60	\$ ::	14.35	11.27	
17.96	15.37	20.21	15.78	10.22	8.10	13.18	14.48	14.23	17.15	11.85	12.15	3.07	13.69	17.62	10.28	15.63	10.79	11.42	11.85	1.8	<u>7</u> .	12.53	12.44	10.85	11.63	13.04	10.73	0.36	. 10.87	13.45	0:30	2.86	1.00	12.46	
0.92	0.51	2.09	4.64	2.05	11.07	11.48	0.88	0.77	11.51	0.93	1.40	12.71	0.67	1.10	11.04	2.05	11.98	5.53	1.15	10.45	11.43	0.97	10.06	12.49	4.58	0.81	3.46	12.50	1.65	12.54	10.05	11.72	10.33	1.45	
10.90	10.84	10.83	10.81	. 10.79	10.70	10.64	10.32	10.21	10.11	10.07	9.95	9.95	68.6	9.87	9.57	9.53	9.52	9.48	9.48	9.47	9.34	9.31	9.24	9.22	9.17	9.12	60.6	9.07	90.6	9.03	8.98	8.87	8.56	8.39	Ω
09.6	6.12	0.31	90.0	9.32	0.09	13.51	0.87	5.55	0.12	2.53	2.29	1.05	4.94	3.82	0.18	0.77	10.61	1.40	1.40	0.09	0.06	0.22	1.03	0.97	9.89	2.44	0.26	0.26	3.21	11.71	17.81	5.97	0.00	9.51	
92	75.0	8	0.0	0.86	0.0	121	80	55.0	0.01	0.25	0.23	0.11	0.50	0.39	0.02	90.0	1.12	0.15	0.15	0.01	0.01	0.02	0.11	0.10	1.08	0.27	0.03	0.03	0.35	1.30	1.98	0.67	0.0	1.13	
040040	4440434	NZ4131	44481513	N20751	44453433	HEBOTA	AA281784	NAGTAB	AA158035	R04504	AA664007	AA625979	AA292226	N21633	AA701652	R46000	N47113	62.206N	AA219060	W16659	AA284072	N50962	H04789	AA705047	AA446462	R96579	R69584	W44766	N26515	AA172056	AA620485	W68220	AA496998	W47364	
102333	75.428	298417	795840	257955	788200	204483	712401	277621	590640	197651	855391	745490	725877	266135	433567	35804	280375	303099	630013	320495	700792	281162	43733	461488	781047	199577	141726	320865	266259	594758	951241	342640	897546	324717	
¥.	3 &	5 12	<u> </u>	3 5	3 5	3	<u>;</u> ;	1 5	? 3	145	. <del>1</del>	7.7	148	149	150		152	153	<del>1</del> 2	155	156	157	58	159	9	161	162	163	2	165	99	167	891	169	

Table 2-12: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line. Genes upregulated 10 fold or more in at least 2 cancer cell lines.

•		7	2
95.07	12.30	11.26	10.67
;		11.14	90.0
	1.30	2.23	11.07
	8.39	8.21	7.27
	3.36	1.49	15.11
	0.40	0.18	2.08
	AA504265	AA058576	AA706968
	825416	380884	451907
	170	171	172

Sequence #	Accession #	Database	Coguenes 52	A.D.C.C.C. 4.0	
Sequence 1	A21185	GENBANK	Sequence 52 Sequence 53	AB033048	GENBANK
Sequence 2	ABC00509	GENBANK	Sequence 54	AB033114	GENBANK
Sequence 3	AB002329	GENBANK	Sequence 55	AD001528	GENBANK
Sequence 4	AB002357	GENBANK	•	AF000986	GENBANK
Sequence 5	AB002366	GENBANK	Sequence 56	AF001893	GENBANK
Sequence 6	AB002387	GENBANK	Sequence 57 Sequence 58	AF002715	GENBANK
Sequence 7	AB004304	GENBANK	•	AF003837	GENBANK
Sequence 8	AB004788	GENBANK	Sequence 59	AF004561	GENBANK
Sequence 9	AB004903	GENBANK	Sequence 60	AF006085	GENBANK
Sequence 10	AB007870	GENBANK	Sequence 61	AF007791	GENBANK
Sequence 11	AB007899	GENBANK	Sequence 62 Sequence 63	AF008442	GENBANK
Sequence 12	AB007930	GENBANK	•	AF010235	GENBANK
Sequence 13	AB007941	GENBANK	Sequence 64	AF013988	GENBANK
Sequence 14	AB007956	GENBANK	Sequence 65	AF016266	GENBANK
Sequence 15	AB007957	GENBANK	Sequence 66	AF016270	GENBANK
Sequence 16	AB007960	GENBANK	Sequence 67	AF019226	GENBANK
Sequence 17	AB009284	GENBANK	Sequence 68	AF023259	GENBANK
Sequence 18	AB011108	GENBANK	Sequence 69 Sequence 70	AF025840	GENBANK
Sequence 19	AB011132	GENBANK	Sequence 70	AF025998	GENBANK
Sequence 20	AB011140	GENBANK	Sequence 72	AF026939	GENBANK
Sequence 21	AB011155	GENBANK	Sequence 72 Sequence 73	AF027824	GENBANK
Sequence 22	AB011159	GENBANK	Sequence 73	AF029890 AF033095	GENBANK
Sequence 23	AB011164	GENBANK	Sequence 75	AF033093 AF034607	GENBANK
Sequence 24	AB011472	GENBANK	Sequence 75	AF035296	GENBANK
Sequence 25	AB012130	GENBANK	Sequence 77	AF035298	GENBANK GENBANK
Sequence 26	AB012910	GENBANK	Sequence 78	AF035309	GENBANK
Sequence 27	AB014521	GENBANK	Sequence 79	AF038404	GENBANK
Sequence 28	AB014589	GENBANK	Sequence 80	AF038957	GENBANK
Sequence 29	AB017007	GENBANK	Sequence 81	AF038960	GENBANK
Sequence 30	AB018268	GENBANK	Sequence 82	AF039703	GENBANK
Sequence 31	AB018305	GENBANK	Sequence 83	AF039918	GENBANK
Sequence 32	AB018327	GENBANK	Sequence 84	AF042331	GENBANK
Sequence 33	AB018344	GENBANK	Sequence 85	AF044588	GENBANK
Sequence 34	AB018351	GENBANK	Sequence 86	AF045167	GENBANK
Sequence 35	AB019568	GENBANK	Sequence 87	AF045184	GENBANK
Sequence 36	AB020637	GENBANK	Sequence 88	AF047020	GENBANK
Sequence 37	AB020669	GENBANK	Sequence 89	AF047472	GENBANK
Sequence 38	<b>AB</b> 0206 <b>80</b>	GENBANK	Sequence 90	AF048977	GENBANK
Sequence 39	AB020682	GENBANK	Sequence 91	AF050171	GENBANK
Sequence 40	AB020692	GENBANK	Sequence 92	AF052124	GENBANK
Sequence 41	AB020723	GENBANK	Sequence 93	AF052164	GENBANK
Sequence 42	AB020981	GENBANK	Sequence 94	AF052182	GENBANK
Sequence 43	AB021288	GENBANK	Sequence 95	AF053470	GENBANK
Sequence 44	AB023153	GENBANK	Sequence 96	AF054179	GENBANK
Sequence 45	AB023206	GENBANK	Sequence 97	AF054187	GENBANK
Sequence 46	AB023227	GENBANK	Sequence 98	AF054838	GENBANK
Sequence 47	AB024704	GENBANK	Sequence 99	AF054990	GENBANK
Sequence 48	AB028624	GENBANK	Sequence 100	AF055012	GENBANK
Sequence 49	AB029000	GENBANK	Sequence 101	AF057160	GENBANK
Sequence 50	AB029004	GENBANK	Sequence 102	AF061258	GENBANK
Sequence 51	AB033020	GENBANK	Sequence 103	AF061938	GENBANK

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Sequence 104	AF067656	GENBANK	Sequence 156	AF155832	GENBANK
Sequence 105	AF067972	GENBANK	Sequence 157	AF159056	GENBANK
Sequence 106	AF068 <b>846</b>	GENBANK	Sequence 158	AF176574	GENBANK
Sequence 107	AF069601	GENBANK	Sequence 159	AF182645	GENBANK
Sequence 108	AF070523	GENBANK	Sequence 160	AF188745	GENBANK
Sequence 109	AF070555	GENBANK	Sequence 161	AF192979	GENBANK
Sequence 110	AF070561	GENBANK	Sequence 162	AJ001306	GENBANK
Sequence 111	AF070655	GENBANK	Sequence 163	AJ004913	GENBANK
Sequence 112	AF070657	GENBANK	Sequence 164	AJ004955	GENBANK
Sequence 113	AF070664	GENBANK	Sequence 165	AJ006470	GENBANK
Sequence 114	AF071202	GENBANK	Sequence 166	AJ006834	GENBANK
Sequence 115	AF071593	GENBANK	Sequence 167	AJ007398	GENBANK
Sequence 116	AF077030	GENBANK	Sequence 168	AJ010442	GENBANK
Sequence 117	AF077034	GENBANK	Sequence 169	AJ010444	GENBANK
Sequence 118	AF077367	GENBANK	Sequence 170	AJ012499	GENBANK
Sequence 119	AF083441	GENBANK	Sequence 171	AJ130733	GENBANK
Sequence 120	AF083470	GENBANK	Sequence 172	AJ224172	GENBANK
Sequence 121	AF085845	GENBANK	Sequence 173	AJ227918	GENBANK
Sequence 121	AF086003	GENBANK	Sequence 174	AJ228139	GENBANK
Sequence 123	AF086218	GENBANK		AJ238095	GENBANK
Sequence 124	AF086336	GENBANK	Sequence 175 Sequence 176	AJ250042	
Sequence 125	AF086431	GENBANK	Sequence 177	AL049227	GENBANK GENBANK
Sequence 126	AF.086495	GENBANK	•	AL049246	
Sequence 127	AF086517	GENBANK	Sequence 178 Sequence 179	AL049240 AL049471	GENBANK GENBANK
Sequence 128	AF086557	GENBANK	Sequence 180	AL049471 AL049799	
•			1		GENBANK
Sequence 129	AF087020	GENBANK	Sequence 181	AL049932	GENBANK
Sequence 130	AF087573	GENBANK	Sequence 182	AL049941	GENBANK
Sequence 131	AF088055	GENBANK	Sequence 183	AL049942	GENBANK
Sequence 132	AF091089 AF093118	GENBANK GENBANK	Sequence 184	AL049969	GENBANK
Sequence 133 Sequence 134	AF100615	GENBANK	Sequence 185	AL049987	GENBANK
•			Sequence 186	AL050041	GENBANK
Sequence 135 Sequence 136	AF100755 AF100756	GENBANK	Sequence 187	AL050141	GENBANK GENBANK
Sequence 137	AF100750 AF101051	GENBANK GENBANK	Sequence 188	AL050161 AL050184	
Sequence 138	AF101031	GENBANK	Sequence 189	AL050198	GENBANK GENBANK
Sequence 139			Sequence 190		
Sequence 140	AF103907 AF106681	GENBANK GENBANK	Sequence 191	AL080172 AL080234	GENBANK GENBANK
Sequence 141	AF110643	GENBANK	Sequence 192	AL096719	GENBANK
Sequence 141	AF112152	GENBANK	Sequence 193 Sequence 194	AL110144	GENBANK
Sequence 142	AF114471	GENBANK	•	AL110183	GENBANK
Sequence 144			Sequence 195		GENBANK
Sequence 145	AF119297	GENBANK	Sequence 196	AL110197	GENBANK
	AF124438	GENBANK	Sequence 197	AL110206	GENBANK
Sequence 146	AF128527	GENBANK	Sequence 198	AL110212	
Sequence 147	AF134159	GENBANK	Sequence 199	AL117237	GENBANK
Sequence 148	AF139461	GENBANK	Sequence 200	AL117427	GENBANK
Sequence 149	AF147380	GENBANK	Sequence 201	AL117429	GENBANK
Sequence 150	AF151103	GENBANK	Sequence 202	AL117505	GENBANK
Sequence 151	AF151840	GENBANK	Sequence 203	AL117534	GENBANK
Sequence 152	AF151903	GENBANK	Sequence 204	AL117554	GENBANK
Sequence 153	AF152306	GENBANK	Sequence 205	AL117576	GENBANK
Sequence 154	AF153191	GENBANK	Sequence 206	AL117609	GENBANK
Sequence 155	AF155099	GENBANK	Sequence 207	AL117643	GENBANK

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Sequence 208	AL133000	GENBANK	Sequence 260	J03209	GENBANK
Sequence 209	D00015	GENBANK	Sequence 261	J03250	GENBANK
Sequence 210	D00017	GENBANK	Sequence 262	J03464	GENBANK
Sequence 211	D13641	GENBANK	Sequence 263	J03779	GENBANK
Sequence 212	D14041	GENBANK	Sequence 264	J03799	GENBANK
Sequence 213	D14696	GENBANK	Sequence 265	J04164	GENBANK
Sequence 214	D16431	GENBANK	Sequence 266	J04621	GENBANK
Sequence 215	D25215	GENBANK	Sequence 267	J04970	GENBANK
Sequence 216	D26488	GENBANK	Sequence 268	J05021	GENBANK
Sequence 217	D28358	GENBANK	Sequence 269	J05032	GENBANK
Sequence 218	D28420	GENBANK	Sequence 270	K01911	GENBANK
Sequence 219	D28482	GENBANK	Sequence 271	K02765	GENBANK
Sequence 220	D29954	GENBANK	Sequence 272	L00160	GENBANK
Sequence 221	D29958	GENBANK	Sequence 273	L01439	GENBANK
Sequence 222	D30037	GENBANK	Sequence 274	L02426	GENBANK
Sequence 223	D31839	GENBANK	Sequence 275	L05425	GENBANK
Sequence 224	D37965	GENBANK	Sequence 276	L06133	GENBANK
Sequence 225	D38551	GENBANK	Sequence 277	L08441	GENBANK
Sequence 226	D38583	GENBANK	Sequence 278	L10678	GENBANK
Sequence 227	D42040	GENBANK	Sequence 279	L10910	GENBANK
Sequence 228	D43950	GENBANK	Sequence 280	L11066	GENBANK
Sequence 229	D50371	GENBANK	Sequence 281	L11932	GENBANK
Sequence 230	D63486	GENBANK	Sequence 282	L13210	GENBANK
Sequence 231	D63874	GENBANK	Sequence 283	L13385	GENBANK
Sequence 232	D80009	GENBANK	Sequence 284	L13773	GENBANK
Sequence 233	D83032	GENBANK	Sequence 285	L13799	GENBANK
Sequence 234	D83197	GENBANK	Sequence 286	L15702	GENBANK
Sequence 235	D83485	GENBANK	Sequence 287	L19161	GENBANK
Sequence 236	D84557	GENBANK	Sequence 288	L19605	GENBANK
Sequence 237	D86228	GENBANK	Sequence 289	L19872	GENBANK
Sequence 238	D86322	GENBANK	Sequence 290	L20941	GENBANK
Sequence 239	D87437	GENBANK	Sequence 291	L22569	GENBANK
Sequence 240	D87444	GENBANK	Sequence 292	L28010	GENBANK
Sequence 241	D87466	GENBANK	Sequence 293	L33930	GENBANK
Sequence 242	D87666	GENBANK	Sequence 294	L36642	GENBANK
Sequence 243	D87667	GENBANK	Sequence 295	L38941	GENBANK
Sequence 244	D87685	GENBANK	Sequence 296	L38961	GENBANK
Sequence 245	D87930	GENBANK	Sequence 297	L39833	GENBANK
Sequence 246	D89053	GENBANK	Sequence 298	L48984	GENBANK
Sequence 247	D89667	GENBANK	Sequence 299	M10119	GENBANK
Sequence 248	D89937	GENBANK	Sequence 300	M11119	GENBANK
Sequence 249	D90359	GENBANK	Sequence 301	M12670	GENBANK
Sequence 250	D90427	GENBANK	Sequence 302	M13231	GENBANK
Sequence 251	E01497	GENBANK	Sequence 303	M13692	GENBANK
Sequence 252	E01956	GENBANK	Sequence 304	M15661	GENBANK
Sequence 253	E01979	GENBANK	Sequence 305	M15885	GENBANK
Sequence 254	E02628	GENBANK	Sequence 306	M15887	GENBANK
Sequence 255	E02822	GENBANK	Sequence 307	M16247	GENBANK
Sequence 256	E02823	GENBANK	Sequence 308	M16342	GENBANK
Sequence 257	E06721	GENBANK	Sequence 309	M16660	GENBANK
Sequence 258	J00194	GENBANK	Sequence 310	M16768	GENBANK
Sequence 259	J03040	GENBANK	Sequence 311	M17324	GENBANK
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Sequence 312	M17517	GENBANK	Sequence 364	U09848	GENBANK
Sequence 313	M17885	GENBANK	Sequence 365	U10439	GENBANK
Sequence 314	M18366	GENBANK	Sequence 366	U13616	GENBANK
Sequence 315	M21302	GENBANK	Sequence 367	U14394	GENBANK
Sequence 316	M22382	GENBANK	Sequence 368	U14510	GENBANK
Sequence 317	M22918	GENBANK	Sequence 369	U16799	GENBANK
Sequence 318	M22920	GENBANK	Sequence 370	U19557	GENBANK
Sequence 319	M24194	GENBANK	Sequence 371	U24105	GENBANK
Sequence 320	M24594	GENBANK	Sequence 372	U24704	GENBANK
Sequence 321	M24902	GENBANK	Sequence 373	U28964	GENBANK
Sequence 322	M25246	GENBANK	Sequence 374	U30521	GENBANK
Sequence 323	M26663	GENBANK	Sequence 375	U34252	GENBANK
Sequence 324	M27332	GENBANK	Sequence 376	U37143	GENBANK
Sequence 325	M27334	GENBANK	Sequence 377	U37283	GENBANK
Sequence 326	M27826	GENBANK	Sequence 378	U37519	
Sequence 327	M28211	GENBANK	Sequence 379	U38654	GENBANK
Sequence 328	M28372	GENBANK	Sequence 380	U39360	GENBANK
Sequence 329	M29064	GENBANK	Sequence 381	U39840	GENBANK
Sequence 330	M29550	GENBANK	Sequence 382	U40671	GENBANK
Sequence 331	M29870	GENBANK	Sequence 383		GENBANK
Sequence 332	M34840	GENBANK	Sequence 384	U41515	GENBANK
Sequence 333	M36341	GENBANK	Sequence 385	U42594	GENBANK
Sequence 334	M37583	GENBANK	Sequence 386	U51134	GENBANK
Sequence 335	M55409	GENBANK	•	U54558	GENBANK
Sequence 336	M55542	GENBANK	Sequence 387	U54831	GENBANK
Sequence 337	M55543	GENBANK	Sequence 388	U56255	GENBANK
Sequence 338	M58485	GENBANK	Sequence 389	U58855	GENBANK
Sequence 339	M59305	GENBANK	Sequence 390	U60975	GENBANK
Sequence 340	M62840	GENBANK	Sequence 391	U62961	GENBANK
Sequence 341	M63838	GENBANK	Sequence 392	U63810	GENBANK
Sequence 342	M69181	GENBANK	Sequence 393	U65928	GENBANK
Sequence 343	M74509	GENBANK	Sequence 394	U66616	GENBANK
Sequence 344	M74558	GENBANK	Sequence 395	U70735	GENBANK
Sequence 345	M74777	GENBANK	Sequence 396	U80034	GENBANK
Sequence 346	M76180	GENBANK	Sequence 397	U80743	GENBANK
Sequence 347	M77830	GENBANK	Sequence 398	U81001	GENBANK
Sequence 348	M80902	GENBANK	Sequence 399	U90028	GENBANK
Sequence 349	M83653	GENBANK	Sequence 400	U90144	GENBANK
Sequence 350	M83738	GENBANK	Sequence 401	U90916	GENBANK .
Sequence 351	M83941	GENBANK	Sequence 402	V00478	GENBANK
Sequence 352	M85164	GENBANK	Sequence 403	X01742	GENBANK
Sequence 353	M87339	GENBANK	Sequence 404	X02422	GENBANK
Sequence 354	M92439		Sequence 405	X02457	GENBANK
Sequence 355	M94654	GENBANK	Sequence 406	X02544	GENBANK
Sequence 356	S70154	GENBANK	Sequence 407	X04098	GENBANK
Sequence 357	S71513	GENBANK	Sequence 408	X04412	GENBANK
Sequence 358		GENBANK	Sequence 409	X04470	GENBANK
Sequence 359	S82240	GENBANK	Sequence 410	X04503	GENBANK
Sequence 360	U02556	GENBANK	Sequence 411	X05276	GENBANK
Sequence 361	U03271	GENBANK	Sequence 412	X05332	GENBANK
Se ence 362	U07681	GENBANK	Sequence 413	X05908	GENBANK
_	U08815	GENBANK	Sequence 414	X07695	GENBANK
Sequence 363	U09564	GENBANK	Sequence 415	X07820	GENBANK

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Sequence 416	X12597	GENBANK	Sequence 468	Z74615	GENBANK
Sequence 417	X13425	GENBANK	Sequence 469	AA005035	dbEST
Sequence 418	X14420	GENBANK	Sequence 470	AA007280	dbEST
Sequence 419	X15187	GENBANK	Sequence 471	AA010019	dbEST
Sequence 420	X15729	GENBANK	Sequence 472	AA010083	dbEST
Sequence 421	X15822	GENBANK	Sequence 473	AA012927	dbEST
Sequence 422	X15880	GE <b>NBANK</b>	Sequence 474	AA016274	dbEST
Sequence 423	X16940	GE <b>NBANK</b>	Sequence 475	AA018892	dbEST
Sequence 424	X52022	GENBANK	Sequence 476	AA019498	dbEST
Sequence 425	X53280	GENBANK	Sequence 477	AA029415	dbEST
Sequence 426	X54941	GENBANK	Sequence 478	AA031483	dbEST
Sequence 427	X55525	GENBANK	Sequence 479	AA031762	dbEST
Sequence 428	X57810	GENBANK	Sequence 480	AA033869	dbEST
Sequence 429	X59618	GE <b>NBANK</b>	Sequence 481	AA034226	dbEST
Sequence 430	X61970	GENBANK	Sequence 482	AA034237	dbEST
Sequence 431	X62744	GENBANK	Sequence 483	AA035742	dbEST
Sequence 432	X63432	GENBANK	Sequence 484	AA037126	dbEST
Sequence 433	X64875	GENBANK	Sequence 485	AA037172	dbEST
Sequence 434	X65550	GENBANK	Sequence 486	AA037554	, dbEST
Sequence 435	X66276	GENBANK	Sequence 487	AA039354	dbEST
Sequence 436	X70326	GENBANK	Sequence 488	AA039504	dbEST
Sequence 437	X71087	GENBANK	Sequence 489	AA040209	dbEST
Sequence 438	X73114	GENBANK	Sequence 490	AA041494	dbEST
Sequence 439	X73902	GENBANK	Sequence 491	AA043431	dbEST
Sequence 440	X74801	GENBANK	Sequence 492	AA044187	dbEST
Sequence 441	X75593	GENBANK	Sequence 493	AA044209	dbEST
Sequence 441	X75861	GENBANK	Sequence 494	AA044416	dbEST
Sequence 443	X76180	GENBANK	Sequence 495	AA044691	dbEST
Sequence 444	X78926	GENBANK	Sequence 496	AA044960	dbEST
Sequence 445	X85372	GENBANK	Sequence 497	AA045584	dbEST
Sequence 446	X87241	GENBANK	Sequence 498	AA047213	dbEST
Sequence 447	X90568	GENBANK	Sequence 499	AA047332	dbEST
Sequence 448	X90870	GENBANK	Sequence 500	AA047704	dbEST
Sequence 449	X91257	GENBANK	Sequence 501	AA053522	dbEST
Sequence 450	X94323	GENBANK	Sequence 502	AA054521	dbEST
Sequence 450	X94754	GENBANK	Sequence 503	AA056281	dbEST
	X95240	GENBANK	Sequence 504	AA056686	dbEST
Sequence 452 Sequence 453	X95677	GENBANK	Sequence 505	AA071030	dbEST
Sequence 454	X99920	GENBANK	Sequence 506	AA074067	dbEST
Sequence 455	Y00052	GENBANK	Sequence 507	AA075008	dbEST
Sequence 456	Y00281	GENBANK	Sequence 508	AA075515	dbEST
Sequence 457	Y00361	GENBANK	Sequence 509	AA075645	dbEST
_			Sequence 510	AA075981	dbEST
Sequence 458	Y00711 Y08915	GENBANK		AA076171	dbEST
Sequence 459		GENBANK	Sequence 511	AA076964	
Sequence 460	Y09267	GENBANK	Sequence 512	AA077283	dbEST dbEST
Sequence 461	Y16241	GENBANK	Sequence 513		dbEST
Sequence 462	Z11338	GENBANK	Sequence 514	AA079706	dbEST
Sequence 463	Z18538	GENBANK	Sequence 515	AA081788	dbEST
Sequence 464	Z23064	GENBANK	Sequence 516	AA081800	dbEST
Sequence 465	Z26317	GENBANK	Sequence 517	AA081973	dbEST
Sequence 466	Z29330	GENBANK	Sequence 518	AA082437	dbEST
Sequence 467	Z36785	GENBANK	Sequence 519	AA083654	dbEST

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Sequence 520	AA086007	dbEST	Soguence E72	A A 400man	
Sequence 521	AA088322	dbEST	Sequence 572	AA186733	dbEST
Sequence 522	AA088850	dbEST	Sequence 573	AA187659	dbEST
Sequence 523	AA092224	dbEST	Sequence 574	AA187817	dbEST
Sequence 524	AA099952	dbEST	Sequence 575	AA188918	dbEST
Sequence 525	AA100987	dbEST	Sequence 576	AA190676	dbEST
Sequence 526	AA101208		Sequence 577	AA194988	dbEST
Sequence 527	AA113403	dbEST	Sequence 578	AA195560	dbEST
Sequence 528	AA115604	dbEST	Sequence 579	AA196978	dbEST
Sequence 529	AA120818	dbEST	Sequence 580	AA203237	dbEST
Sequence 530	AA121471	dbEST	Sequence 581	AA205870	dbEST
Sequence 531	AA121923	dbEST	Sequence 582	AA211498	dbEST
Sequence 532	AA125809	dbEST	Sequence 583	AA213548	dbEST
Sequence 533		dbEST	Sequence 584	AA215800	dbEST
Sequence 534	AA125989	dbEST	Sequence 585	AA218858	dbEST
Sequence 535	AA126128	dbEST	Sequence 586	AA223425	dbEST
Sequence 536	AA127674	dbEST	Sequence 587	AA224230	dbEST
Sequence 537	AA128306	dbEST	Sequence 588	AA225025	dbEST
	AA128391	dbEST	Sequence 589	AA229993	dbEST
Sequence 538	AA129501	dbEST	Sequence 590	AA232738	dbEST
Sequence 539	AA129909	dbEST	Sequence 591	AA233867	dbEST
Sequence 540	AA132798	dbEST	Sequence 592	AA234149	dbEST
Sequence 541	AA132956	dbEST	Sequence 593	AA234464	dbEST
Sequence 542	AA133382	dbEST	Sequence 594	AA234646	dbEST
Sequence 543	AA133503	dbEST	Sequence 595	AA234667	dbEST
Sequence 544	AA134032	dbEST	Sequence 596	AA247638	dbEST
Sequence 545	AA134778	dbEST	Sequence 597	AA250846	dbEST
Sequence 546	AA135049	dbEST	Sequence 598	AA250854	dbEST
Sequence 547	AA135104	dbEST	Sequence 599	AA253248	dbEST
Sequence 548	AA136794	dbEST	Sequence 600	AA255577	dbEST
Sequence 549	AA136830	dbEST	Sequence 601	AA256290	dbEST
Sequence 550	AA148002	dbEST	Sequence 602	AA256297	dbEST
Sequence 551	AA149624	dbEST	Sequence 603	AA256591	dbEST
Sequence 552	AA150928	dbEST	Sequence 604	AA258003	dbEST
Sequence 553	AA156040	dbEST	Sequence 605	AA261968	dbEST
Sequence 554	AA159847	dbEST	Sequence 606	AA262683	dbEST
Sequence 555	AA160002	dbEST	Sequence 607	AA262783	dbEST
Sequence 556	AA161003	dbEST	Sequence 608	AA278357.	dbEST
Sequence 557	AA164473	dbEST	Sequence 609	AA278785 -	dbEST
Sequence 558	AA165243	dbEST	Sequence 610	AA278956	dbEST
Sequence 559	AA165632	dbEST	Sequence 611	AA279145	dbEST
Sequence 560	AA165639	dbEST	Sequence 612	AA279262	dbEST
Sequence 561	AA166907	dbEST	Sequence 613	AA282956	dbEST
Sequence 562	AA169276	dbEST	Sequence 614	AA283112	dbEST
Sequence 563	AA169638	dbEST	Sequence 615	AA285043	dbEST
Sequence 564	AA171516	dbEST	Sequence 616	AA291436	dbEST
Sequence 565	AA171872	dbEST	Sequence 617	AA292132	dbEST
Sequence 566	AA179105	dbEST	Sequence 618	AA292281	dbEST
Sequence 567	AA180270	dbEST	Sequence 619	AA293027	dbEST
Sequence 568	AA181526	dbEST	Sequence 620	AA295182	dbEST
Sequence 569	AA181580	dbEST	Sequence 621	AA295996	dbEST
Sequence 570	AA182409	dbEST	Sequence 622	AA297432	dbEST
Sequence 571	AA186538	dbEST	Sequence 623	AA298085	
		,	ocquence 023	~~~30000	dbES <b>T</b>

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Sequence 624	AA298773	dbEST	Sequence 676	AA417324	dbEST
Sequence 625	AA298794	dbEST	Sequence 677	AA418061	dbEST
Sequence 626	AA300440	dbEST	Sequence 678	AA418080	dbEST
Sequence 627	AA303257	dbEST	Sequence 679	AA418681	dbEST
Sequence 628	AA304979	dbEST	Sequence 680	AA419263	dbEST
Sequence 629	AA305319	db <b>EST</b>	Sequence 681	AA421682	dbEST
Sequence 630	AA305494	dbEST	Sequence 682	AA423943	dbEST
Sequence 631	AA306772	dbEST	Sequence 683	AA425218	dbEST
Sequence 632	AA307286	dbEST	Sequence 684	AA425619	dbEST
Sequence 633	AA307539	dbEST	Sequence 685	AA428329	dbEST
Sequence 634	AA307857	dbEST	Sequence 686	AA429721	dbEST
Sequence 635	AA309002	dbEST	Sequence 687	AA430381	dbEST
Sequence 636	AA309127	dbEST	Sequence 688	AA430400	dbEST
Sequence 637	AA310469	dbEST	Sequence 689	AA431226	dbEST
Sequence 638	AA311028	dbEST	Sequence 690	AA432075	dbEST
Sequence 639	AA312078	dbEST	Sequence 691	AA435887	dbEST
Sequence 640	AA312962	dbEST	Sequence 692	AA436765	dbEST
Sequence 641	AA313164	dbEST	Sequence 693	AA443154	dbEST
Sequence 642	AA313653	dbEST	Sequence 694	AA443806	dbEST
Sequence 643	AA313692	dbEST	Sequence 695	AA443870	dbEST
Sequence 644	AA313922	dbEST	Sequence 696	AA447116	dbEST
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Sequence 648	AA319601	dbEST	Sequence 700	AA453486	dbEST
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Sequence 651	AA334754	dbEST	Sequence 703	AA456737	dbEST
Sequence 652	AA340635	dbEST	Sequence 704	AA458921	dbEST
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Sequence 658	AA357574	dbEST	Sequence 710	AA468335	dbEST
Sequence 659	AA360977	dbEST	Sequence 711	AA468839	dbEST
Sequence 660	AA361225	dbEST '	Sequence 712	AA476392	dbEST
Sequence 661	AA363049	dbEST	Sequence 713	AA477288	dbEST
Sequence 662	AA371483	dbEST	Sequence 714	AA477856	dbEST
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Sequence 666	AA398214	dbEST	Sequence 718	AA482430	dbEST
Sequence 667	AA398463	dbEST	Sequence 719	AA482434	dbEST
Sequence 668	AA398674	dbEST	Sequence 720	AA483635	dbEST
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Sequence 672	AA401208	dbEST	Sequence 724	AA489032	dbEST
Sequence 673	AA401285	dbEST	Sequence 725	AA490635	dbEST
Sequence 674	AA410580	dbEST	Sequence 726	AA492272	dbEST
Sequence 675	AA411736	dbEST	Sequence 727	AA492280	dbEST
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Sequence 728	AA492392	dbEST	Sequence 780	AA586969	dbEST
Sequence 729	AA493512	dbEST	Sequence 781	AA587630	dbEST
Sequence 730	AA502979	dbEST	Sequence 782	AA594282	dbEST
Sequence 731	AA503330	dbEST	Sequence 783	AA594780	dbEST
Sequence 732	AA505400	dbEST	Sequence 784	AA594950	dbEST
Sequence 733	AA507314	dbEST	Sequence 785	AA599424	dbEST
Sequence 734	AA508804	dbEST	Sequence 786	AA599533	dbEST
Sequence 735	AA514933	dbEST	Sequence 787	AA602361	dbEST
Sequence 736	AA522686	dbEST	Sequence 788	AA602957	dbEST
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Sequence 746	AA531255	dbEST	Sequence 798	^A631178	dbEST
Sequence 747	AA531563	dbEST	Sequence 799	AA631206	dbEST
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Sequence 753	AA548098	dbEST	Sequence 805	AA653508	dbEST
Sequence 754	AA548244	dbEST	Sequence 806	AA654995	dbEST
Sequence 755	AA548600	dbEST	Sequence 807	AA657463	dbEST
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Sequence 758	AA554010	dbEST	Sequence 810	AA669976	dbEST
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Sequence 760	AA554718	dbEST	Sequence 812	AA687495	dbEST
Sequence 761	AA555227	dbEST	Sequence 813	AA702422	dbEST
Sequence 762	AA558280	dbEST	Sequence 814	AA704005	dbEST
Sequence 763	AA564843	dbEST	Sequence 815	AA705061	dbEST
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Sequence 766	AA572950	dbEST	Sequence 818	AA729119	dbEST
Sequence 767	AA573742	dbEST	Sequence 819	AA740151	dbEST
Sequence 768	AA574397	dbEST	Sequence 820	AA744239	dbEST
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Sequence 775	AA583431	dbEST	Sequence 827	AA758592	dbEST
Sequence 776	AA583567	dbEST	Sequence 828	AA767439	dbEST
Sequence 777	AA583773	dbEST	Sequence 829	AA769127	dbEST
Sequence 778	AA583899	dbEST	Sequence 830	AA769937	dbEST
Sequence 779	AA583997	dbEST	Sequence 831	AA774523	dbEST
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Sequence 832	AA774750	dbEST	Sequence 884	AA988468	dbEST
Sequence 833	AA777489	dbEST	Sequence 885	AA992583	dbEST
Sequence 834	AA779728	dbEST	Sequence 886	AA992596	dbEST
Sequence 835	AA779949	dbEST	Sequence 887	AA993550	dbEST
Sequence 836	AA782592	dbEST	Sequence 888	AF001542	dbEST
Sequence 837	AA806040	dbEST	Sequence 889	AF010244	dbEST
Sequence 838	AA814414	dbEST	Sequence 890	AF156972	dbEST
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Sequence 840	AA827331	dbEST	Sequence 892	AI016113	dbEST
Sequence 841	AA834102	dbEST	Sequence 893	AI016417	dbEST
Sequence 842	AA834946	dbEST	Sequence 894	AI017236	dbEST
Sequence 843	AA845374	dbEST	Sequence 895	AI018468	dbEST
Sequence 844	AA847210	dbEST	Sequence 896	AI018578	dbEST
Sequence 845	AA854319	dbEST	Sequence 897	AI021925	dbEST
Sequence 846	AA858389	dbEST	Sequence 898	AI022353	dbEST
Sequence 847	AA858396	dbEST	Sequence 899	AI026819	dbEST
Sequence 848	AA860287	dbEST	Sequence 900	AI027232	dbEST
Sequence 849	AA861428	dbEST	Sequence 901	AI031819	dbEST
Sequence 850	AA861631	dtEST	Sequence 902	AI033904	dbEST
Sequence 851	AA872729	dbEST	Sequence 903	AI038986	dbEST
Sequence 852	AA873110	dbEST	Sequence 904	Al041820	dbEST
Sequence 853	AA877864	dbEST	Sequence 905	AI042290	dbEST
Sequence 854	AA878781	dbEST	Sequence 906	AI051987	dbEST
Sequence 855	AA883580	dbEST	Sequence 907	AI052124	dbEST
Sequence 856	AA884922	dbEST	Sequence 908	AI052525	dbEST
Sequence 857	AA888121	dbEST	Sequence 909	AI064691	dbEST
Sequence 858	AA888589	dbEST	Sequence 910	AI074713	dbEST
Sequence 859	AA890660	dbEST	Sequence 911	AI075189	dbEST
Sequence 860	AA906652	dbEST	Sequence 912	AI075211	dbEST
Sequence 861	AA910369	dbEST	Sequence 913	AI079329	dbEST
Sequence 862	AA911154	dbEST	Sequence 914	AI080377	dbEST
Sequence 863	AA911317	dbEST	Sequence 915	AI082168	dbEST
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Sequence 868	AA928420	dbEST	Sequence 920	AI091425	dbEST
Sequence 869	AA934879	dbEST	Sequence 921	AI092428	dbEST
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Sequence 871	AA936187	dbEST	Sequence 923	AI093553	dbEST
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Sequence 876	AA962550	dbEST	Sequence 928	Al114467	dbEST
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Sequence 882	AA984043	dbEST	Sequence 934	AI133387	dbEST
Sequence 883	AA984690	dbEST	Sequence 935	AI141847	dbEST
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Sequence 948	AI198416	dbEST	Sequence 1000	Al361377	dbEST
Sequence 949	Al200830	dbEST	Sequence 1001	Al366459	dbEST
Sequence 950	AI207570	dbEST	Sequence 1002	Al366549	dbEST
Sequence 951	AI207628	dbEST	Sequence 1003	Al368048	dbEST
Sequence 952	Al214054	dbEST	Sequence 1004	AI368114	dbEST
Sequence 953	Al215841	dbEST	Sequence 1005	Al371832	dbEST
Sequence 954	Al216971	dbEST	Sequence 1006	Al374643	dbEST
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Sequence 959	Al219116	dbEST	Sequence 1011	AI379402	dbEST
Sequence 960	Al220151	dbEST	Sequence 1012	Al381351	dbEST
Sequence 961	Al221894	dbEST	Sequence 1013	AI382980	dbEST
Sequence 962	Al223085	dbEST	Sequence 1014	A1392607	dbEST
Sequence 963	Al241212	dbEST	Sequence 1015	Al399717	dbEST
Sequence 964	Al242761	dbEST	Sequence 1016	AI400752	dbEST
Sequence 965	Al243343	dbEST	Sequence 1017	AI421761	dbEST
Sequence 966	Al249866	dbEST	Sequence 1018	AI422243	dbEST
Sequence 967	Al253330	dbEST	Sequence 1019	Al431675	dbEST
Sequence 968	Al253381	dbEST	Sequence 1020	AI431869	dbEST
Sequence 969	Al253436	dbEST	Sequence 1021	Al432969	dbEST
Sequence 970	Al261899	dbEST	Sequence 1022	Ai433717	dbEST
Sequence 971	Al267162	dbEST	Sequence 1023	Al439248	dbEST
Sequence 972	Al267185	dbEST	Sequence 1024	Al439762	dbEST
Sequence 973	AI267282	dbEST	Sequence 1025	Al445471	dbEST
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Sequence 978	Al267502	dbEST	Sequence 1030	Al474750	dbEST
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Sequence 985	Al278838	dbEST	Sequence 1037	AI557182	dbEST
Sequence 986	Al280637	dbEST	Sequence 1038	AI557225	dbEST
Sequence 987	Al281509	dbEST	Sequence 1039	AI557495	dbEST

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Sequence 1040	AI557599	dbEST	Sequence 1092	AI753269	dbEST
Sequence 1041	AI557626	dbEST	Sequence 1093	AI754437	dbEST
Sequence 1042	A1559680	dbEST	Sequence 1094	A1755085	dbEST
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Sequence 1048	AI573140	dbEST	Sequence 1100	Al800896	dbEST
Sequence 1049	AI573212	dbEST	Sequence 1101	Al804346	dbEST
Sequence 1050	AI582365	dbEST	Sequence 1102	AI804983	dbEST
Sequence 1051	AI587300	dbEST	Sequence 1103	AI810626	dbEST
Sequence 1052	AI588866	dbEST	Sequence 1104	AI811498	dbEST
Sequence 1053	AI589285	dbEST	Sequence 1105	AI814148	dbEST
Sequence 1054	AI589756	dbEST	Sequence 1106	Al815829	dbEST
Sequence 1055	Al613093	dbEST	Sequence 1107	AI819225	dbEST
Sequence 1056	AI623176	dbEST	Sequence 1108	AI819564	dbEST
Sequence 1057	A1623804	dbEST	Sequence 1109	AI819701	dbEST
Sequence 1058	Al624249	dbEST	Sequence 1110	AI859619	dbEST
Sequence 1059	Al624441	dbEST	Sequence 1111	AI869280	dbEST
Sequence 1060	AI628885	dbEST	Sequence 1112	AI879992	dbEST
Sequence 1061	AI631745	dbEST	Sequence 1113	Al890387	dbEST
Sequence 1062	AI634702	dbEST	Sequence 1114	AI890616	dbEST
Sequence 1063	AI635096	dbEST	Sequence 1115	AI911059	dbEST
Sequence 1064	AI654284	dbEST	Sequence 1116	Al914030	dbEST
Sequence 1065	AI654454	dbEST	Sequence 1117	Al914128	dbEST
Sequence 1066	AI655189	dbEST	Sequence 1118	AI916584	dbEST
Sequence 1067	AI656323	dbEST	Sequence 1119	AI922760	dbEST
Sequence 1068	AI658689	dbEST	Sequence 1120	AI923574	dbEST
Sequence 1069	AI680218	dbEST	Sequence 1121	AI927301	dbEST
Sequence 1070	A1680242	dbEST	Sequence 1122	AI936548	dbEST
Sequence 1071	A1682287	dbEST	Sequence 1123	Al952294	dbEST
Sequence 1072	A1683094	dbEST	Sequence 1124	AI955769	dbEST
Sequence 1073	AI683431	dbEST	Sequence 1125	Al984541	dbEST
Sequence 1074	AI684157	dbEST	Sequence 1126	AI984763	dbEST
Sequence 1075	Al688187	dbEST	Sequence 1127	AJ243229	dbEST
Sequence 1076	Al6 <b>89722</b>	dbEST	Sequence 1128	AL035998	dbEST
Sequence 1077	AI693493	dbEST	Sequence 1129	AL036354	dbEST
Sequence 1078	Al69 <b>784</b> 9	dbEST	Sequence 1130	AL037724	dbEST
Sequence 1079	Al698198	dbEST	Sequence 1131	AL039293	dbEST
Sequence 1080	A1703405	dbEST	Sequence 1132	AL040084	dbEST
Sequence 1081	A1720403	dbEST	Sequence 1133	AL040993	dbEST
Sequence 1082	AI733862	dbEST	Sequence 1134	AL042815	db <b>EST</b>
Sequence 1083	AI740534	dbEST	Sequence 1135	AL043346	dbEST
Sequence 1084	AI740792	dbEST	Sequence 1136	AL045297	dbEST
Sequence 1085	A1740796	dbEST	Sequence 1137	AL110357	dbEST
Sequence 1086	A1740900	dbEST	Sequence 1138	AL119728	dbEST
Sequence 1087	A1743852	dbEST	Sequence 1139	AW003247	dbEST
Sequence 1088	A1750535	dbEST	Sequence 1140	AW003967	dbEST
Sequence 1089	AI750567	dbEST	Sequence 1141	AW009562	dbEST
Sequence 1090	AI750846	dbEST	Sequence 1142	AW015055	dbEST
Sequence 1091	A1751565	dbEST	Sequence 1143	AW025074	dbEST

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Sequence 1144		dbEST	Sequence 1196	N99698	dbEST
Sequence 1145	AW027586	dbEST	Sequence 1197	R13761	dbEST
Sequence 1146	AW044029	dbEST	Sequence 1198	R33779	dbEST
Sequence 1147		dbEST	Sequence 1199	R48977	dbEST
Sequence 1148	AW068491	dbEST	Sequence 1200	R50104	dbEST
Sequence 1149	AW090544	dbEST	Sequence 1201	R61734	dbEST
Sequence 1150	AW104641	dbEST	Sequence 1202	R64693	dbEST
Sequence 1151	AW118812	dbEST	Sequence 1203	R6 <b>746</b> 6	dbEST
Sequence 1152	AW157303	dbEST	Sequence 1204	R70639	dbEST
Sequence 1153	AW170193	dbEST	Sequence 1205	R77186	dbEST
Sequence 1154	C04376	dbEST	Sequence 1206	R82429	dbEST
Sequence 1155	F07019	dbEST	Sequence 1207	R96147	dbEST
Sequence 1156	F07647	dbEST	Sequence 1208	R99043	dbEST
Sequence 1157	H02308	dbEST	Sequence 1209	T78419	dbEST
Sequence 1158	H03785	dbEST	Sequence 1210	T88702	dbEST
Sequence 1159	H08511	dbEST	Sequence 1211	W05463	dbEST
Sequence 1160	H08542	dbEST	Sequence 1212	W15477	dbEST
Sequence 1161	H08959	dbEST	Sequence 1213	W56388	dbEST
Sequence 1162	H10107	dbEST	Sequence 1214	W87522	dbEST
Sequence 1163	H10555	dbEST	Sequence 1215	W89057	dbEST
Sequence 1164	H12229	dbEST	Sequence 1216	Z25142	dbEST
Sequence 1165	H12519	dbEST	Sequence 1217	Z36731	dbEST
Sequence 1166	H14985	dbEST	Sequence 1218	Q37741	NUCPATENT
Sequence 1167	H22734	dbEST	Sequence 1219	T19093	NUCPATENT
Sequence 1168	H23157	dbEST	Sequence 1220	T59274	NUCPATENT
Sequence 1169	H26678	dbEST	Sequence 1221	V43261	NUCPATENT
Sequence 1170	H29664	dbEST	Sequence 1222	V43613	NUCPATENT
Sequence 1171	H38914	dbEST	Sequence 1223	V59617	NUCPATENT
Sequence 1172	H54680	dbEST	Sequence 1224	V62427	NUCPATENT
Sequence 1173	H57585	dbEST	Sequence 1225	V62428	NUCPATENT
Sequence 1174	H57648	dbEST	Sequence 1226	V63637	NUCPATENT
Sequence 1175	H61886	dbEST	Sequence 1227	V74172	NUCPATENT
Sequence 1176	H64719	dbEST	Sequence 1228	V84428	NUCPATENT
Sequence 1177	H73335	dbEST	Sequence 1229	X02974	NUCPATENT
Sequence 1178	H83784	dbEST	Sequence 1230	X03841	NUCPATENT
Sequence 1179	H8 <b>4729</b>	dbEST	Sequence 1231	X04408	NUCPATENT
Sequence 1180	H92106	dbEST	Sequence 1232	X22111	NUCPATENT
Sequence 1181	H94927	dbEST	Sequence 1233	X27262	NUCPATENT
Sequence 1182	H99611	dbEST	Sequence 1234	X27329	NUCPATENT
Sequence 1183	M62015	dbEST	Sequence 1235	X30167	NUCPATENT
Sequence 1184	N25751	dbEST	Sequence 1236	X37486	NUCPATENT
Sequence 1185	N28308	dbEST	Sequence 1237	X976 <b>77</b>	NUCPATENT
Sequence 1186	N36968	dbEST	Sequence 1238	Z18356	NUCPATENT
Sequence 1187	N42196	dbEST	Sequence 1239	Z24611	NUCPATENT
Sequence 1188	N42772	dbEST	Sequence 1240	AC31093	PREPATNUC
Sequence 1189	N63991	dbEST	•		
Sequence 1190	N77277	dbEST	Sequence #	Accession #	Database
Sequence 1191	N78279	dbEST	Sequence 1241	A18657	GENBANK
Sequence 1192	N81205	dbEST	Sequence 1242	AB000095	GENBANK
Sequence 1193	N91096	dbEST	Sequence 1243	AB000220	GENBANK
Sequence 1194	N94855	dbEST	Sequence 1244	AB002366	GENBANK
Sequence 1195	N99553	dbEST	Sequence 1245	AB002387	GENBANK
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Sequence 1246		GENBANK	Sequence 1298	AF021819	CENIDANIZ
Sequence 1247		GENBANK	Sequence 1299	AF026166	GENBANK
Sequence 1248	AB007883	GENBANK	Sequence 1300	AF026291	GENBANK
Sequence 1249	AB008109	GENBANK	Sequence 1301	AF027824	GENBANK
Sequence 1250	AB009285	GENBANK	Sequence 1302	AF031647	GENBANK
Sequence 1251	AB010882	GENBANK	Sequence 1303	AF031647 AF032885	GENBANK
Sequence 1252	AB011175	GENBANK	Sequence 1304	AF032865 AF034759	GENBANK
Sequence 1253	AB013897	GENBANK	Sequence 1305	AF035286	GENBANK
Sequence 1254	AB014511	GENBANK	Sequence 1306	AF035537	GENBANK
Sequence 1255	AB014514	GENBANK	Sequence 1307		GENBANK
Sequence 1256	AB014525	GENBANK	Sequence 1308	AF035839 AF038451	GENBANK
Sequence 1257	AB014531	GENBANK	Sequence 1309		GENBANK
Sequence 1258	AB014536	GENBANK	Sequence 1310	AF038963	GENBANK
Sequence 1259	AB014563	GENBANK	Sequence 1311	AF039918	GENBANK
Sequence 1260	AB014569	GENBANK		AF041260	GENBANK
Sequence 1261	AB014610	GENBANK	Sequence 1312	AF042346	GENBANK
Sequence 1262	AB018257	GENBANK	Sequence 1313	AF045167	GENBANK
Sequence 1263	AB018304	GENBANK	Sequence 1314	AF047020	GENBANK
Sequence 1264	AB018344	GENBANK	Sequence 1315	AF047185	GENBANK
Sequence 1265	AB019002	GENBANK	Sequence 1316	AF047440	GENBANK
Sequence 1266	AB019563	GENBANK	Sequence 1317	AF050637	GENBANK
Sequence 1267	AB019568	GENBANK	Sequence 1318	AF050638	GENBANK
Sequence 1268	AB020637	GENBANK	Sequence 1319	AF051941	GENBANK
Sequence 1269	AB020686	GENBANK	Sequence 1320	AF052138	GENBANK
Sequence 1270	AB020692	GENBANK	Sequence 1321	AF052153	GENBANK
Sequence 1271	AB020707	GENBANK	Sequence 1322	AF052159	GENBANK
Sequence 1272	AB021179	GENBANK	Sequence 1323	AF052183	GENBANK
Sequence 1273	AB021288	GENBANK	Sequence 1324	AF052642	GENBANK
Sequence 1274	AB022663	GENBANK	Sequence 1325	AF054284	GENBANK
Sequence 1275	AB023145	GENBANK	Sequence 1326	AF054838	GENBANK
Sequence 1276	AB028969	GENBANK	Sequence 1327	AF061258	GENBANK
Sequence 1277	AB029036	GENBANK	Sequence 1328	AF061737	GENBANK
Sequence 1278	AB032969	GENBANK	Sequence 1329	AF065388	GENBANK
Sequence 1279	AB032990	GENBANK	Sequence 1330	AF065391	GENBANK
Sequence 1280	AB033029	GENBANK	Sequence 1331	AF068180	GENBANK
Sequence 1281	AB033069	GENBANK	Sequence 1332	AF068235	GENBANK
Sequence 1282	AB033078	GENBANK	Sequence 1333	AF068754	GENBANK
Sequence 1283	AF001893	GENBANK	Sequence 1334	AF069762	GENBANK
Sequence 1284	AF004162	GENBANK	Sequence 1335	AF070523	GENBANK
Sequence 1285	AF004327	GENBANK	Sequence 1336	AF070553	GENBANK
Sequence 1286	AF004813	GENBANK	Sequence 1337	AF070561	GENBANK
Sequence 1287	AF006514		Sequence 1338	AF070562	GENBANK
Sequence 1288	AF007135	GENBANK	Sequence 1339	AF070595	GENBANK
Sequence 1289	AF007216	GENBANK	Sequence 1340	AF070626	GENBANK
Sequence 1290	AF007791	GENBANK	Sequence 1341	AF070650	GENBANK
Sequence 1291	AF008915	GENBANK	Sequence 1342	AF070658	GENBANK
Sequence 1292	AF010233	GENBANK	Sequence 1343	AF070668	GENBANK
Sequence 1293		GENBANK	Sequence 1344	AF071202	GEN <b>BANK</b>
Sequence 1294	AF015040	GENBANK	Sequence 1345	AF073298	GENBANK
Sequence 1295	AF016369	GENBANK	Sequence 1346	AF073475	GENBANK
Sequence 1296	AF020038	GENBANK	Sequence 1347	AF077029	GENBANK
Sequence 1297	AF020202	GENBANK	Sequence 1348	AF077032	GENBANK
ocquerice 1297	AF020736	GENBANK	Sequence 1349	AF077045	GENBANK

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Sequence 13		GENBANK	Sequence 1402	AF191018	GENBANK
Sequence 13		GENBANK	Sequence 1403	AF191020	GENBANK
Sequence 13		GENBANK	Sequence 1404	AF193795	GENBANK
Sequence 13		GENBANK	Sequence 1405	AF195417	GENBANK
Sequence 13		GENBANK	Sequence 1406	AF201077	GENBANK
Sequence 13		GENBANK	Sequence 1407	AJ004913	GENBANK
Sequence 13		GENBANK	Sequence 1408	AJ006834	GENBANK
Sequence 13		GENBANK	Sequence 1409	AJ007398	GENBANK
Sequence 13		GENBANK	Sequence 1410	AJ007714	GENBANK
Sequence 13	59 AF085850	GENBANK	Sequence 1411	AJ010071	GENBANK
Sequence 136	60 AF086172	GENBANK	Sequence 1412	AJ130733	GENBANK
Sequence 136	61 AF086336	GENBANK	Sequence 1413	AJ224172	GENBANK
Sequence 136	52 AF086351	GENBANK	Sequence 1414	AJ224819	GENBANK
Sequence 136	33 AF086418	GENBANK	Sequence 1415	AJ228139	GENBANK
Sequence 136	64 AF086495	GENBANK	Sequence 1416	AL035304	GENBANK
Sequence 136	S5 AF086557	GENBANK	Sequence 1417	AL049233	GENBANK
Sequence 136	66 AF090327	GENBANK	Sequence 1418	AL049252	GENBANK
Sequence 136	67 AF091071	GENBANK	Sequence 1419	AL049325	GENBANK
Sequence 136		GENBANK	Sequence 1420	AL049447	GENBANK
Sequence 136	9 AF095687	GENBANK	Sequence 1421	AL049969	GENBANK
Sequence 137	'0 AF098951	GENBANK	Sequence 1422	AL049987	GENBANK
Sequence 137	1 AF100741	GENBANK	Sequence 1423	AL050018	GENBANK
Sequence 137	<sup>7</sup> 2 AF100756	GENBANK	Sequence 1424	AL050041	GENBANK
Sequence 137	'3 AF102265	GENBANK	Sequence 1425	AL050089	GENBANK
Sequence 137		GENBANK	Sequence 1426	AL050197	GENBANK
Sequence 137	'5 AF103907	GENBANK	Sequence 1427	AL050198	GENBANK
Sequence 137	'6 AF111713	GENBANK	Sequence 1428	AL050255	GENBANK
Sequence 137		GENBANK	Sequence 1429	AL050265	GENBANK
Sequence 137		GENBANK	Sequence 1430	AL050272	GENBANK
Sequence 137		GENBANK	Sequence 1431	AL050290	GENBANK
Sequence 138		GENBANK	Sequence 1432-	AL050373	GENBANK
Sequence 138		GENBANK	Se nce 1433	AL079298	GENBANK
Sequence 138		GENBANK	Se: nce 1434	AL080063	GENBANK
Sequence 138		GENBANK	Sequence 1435	AL080089	GENBANK
Sequence 138	_	GENBANK	Sequence 1436	AL080135	GENBANK
Sequence 138	_	GENBANK	Sequence 1437	AL080144	GENBANK
Sequence 138		GENBANK	Sequence 1438	AL080224	GENBANK
Sequence 138		GENBANK	Sequence 1439	AL110183	GENBANK
Sequence 138		GENBANK	Sequence 1440	AL117237	GENBANK
Sequence 138		GENBANK	Sequence 1441	AL117412	GENBANK
Sequence 139		GENBANK	Sequence 1442	AL117423	GENBANK
Sequence 139		GENBANK	Sequence 1443	AL117429	GENBANK
Sequence 139		GENBANK	Sequence 1444	AL117516	GENBANK '
Sequence 139		GENBANK	Sequence 1445	AL117543	GENBANK
Sequence 139		GENBANK	Sequence 1446	AL117554	GENBANK
Sequence 139		GENBANK	Sequence 1447	AL117584	GENBANK
Sequence 139		GENBANK	Sequence 1448	AL117595	GENBANK
Sequence 139		GENBANK	Sequence 1449	AL122079	GENBANK
Sequence 139		GENBANK	Sequence 1450	D00860	GENBANK
Sequence 1399		GENBANK	Sequence 1451	D10040	GENBANK
Sequence 1400		GENBANK	Sequence 1452	D13119	GENBANK
Sequence 140	1 AF188745	GENBANK	Sequence 1453	D13286	GENBANK

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Sequence 1454	D13643	GENBANK	Sequence 1506	D89077	GENBANK
Sequence 1455	D13645	GENBANK	Sequence 1507	D89667	GENBANK
Sequence 1456	D13748	GENBANK	Sequence 1508	E01888	GENBANK
Sequence 1457	D13757	GENBANK	Sequence 1509	E01956	GENBANK
Sequence 1458	D13866	GENBANK	Sequence 1510	E02628	GENBANK
Sequence 1459	D13900	GENBANK	Sequence 1511	E03569	GENBANK
Sequence 1460	D14659	GENBANK	Sequence 1512	E03814	GENBANK
Sequence 1461	D14697	GENBANK	Sequence 1513	E05957	GENBANK
Sequence 1462	D14812	GENBANK	Sequence 1514	E08515	GENBANK
Sequence 1463	D15057	GENBANK	<b>Sequence 151</b> 5	J02959	GENBANK
Sequence 1464	D16217	GENBANK	Sequence 1516	J03248	GENBANK
Sequence 1465	D16480	GENBANK	Sequence 1517	J03799	GENBANK
Sequence 1466	D17039	GENBANK	Sequence 1518	J04080	GENBANK
Sequence 1467	D17652	GENBANK	Sequence 1519	J04973	GENBANK
Sequence 1468	D21209	GENBANK	Sequence 1520	J05021	GENBANK
Sequence 1469	D21262	GENBANK	Sequence 1521	J05192	GENBANK
Sequence 1470	D23662	GENBANK	Sequence 1522	K01911	GENBANK
Sequence 1471	D25542	GENBANK	Sequence 1523	K03002	GENBANK
Sequence 1472	D28473	GENBANK	Sequence 1524	l 02547	GENBANK
Sequence 1473	D29954	GENBANK	Sequence 1525	L06070	GENBANK
Sequence 1474	D30655	GENBANK	Sequence 1526	L07033	GENBANK
Sequence 1475	D31767	GENBANK	Sequence 1527	L07261	GENBANK
Sequence 1476	D31883	GENBANK	Sequence 1528	L07515	GENBANK
Sequence 1477	D31884	GENBANK	Sequence 1529	L08441	GENBANK
Sequence 1478	D31891	GENBANK	Sequence 1530	L11284	GENBANK
Sequence 1479	D37991	GENBANK	Sequence 1531	L11667	GENBANK
Sequence 1480	D38047	GENBANK	Sequence 1532	L12168	GENBANK
Sequence 1481	D38293	GENBANK	Sequence 1533	L13806	GENBANK
Sequence 1482	D38491	GENBANK	Sequence 1534	L14837	GENBANK
Sequence 1483	D38521	GENBANK	Sequence 1535	L14848	GENBANK
Sequence 1484	D38524	GENBANK	Sequence 1536	L16558	GENBANK
Sequence 1485	D43948	GENBANK	Sequence 1537	L19184	GENBANK
Sequence 1486	D43951	GENBANK	Sequence 1538	L19437	GENBANK
Sequence 1487	D44466	GENBANK	Sequence 1539	L20688	GENBANK
Sequence 1488	D50371	GENBANK	Sequence 1540	L31801	GENBANK
Sequence 1489	D50372	GENBANK	Sequence 1541	L34840	GENBANK
Sequence 1490	D50525	GENBANK	Sequence 1542	L37080	GENBANK
Sequence 1491	D50645	GENBANK	Sequence 1543	L38486	GENBANK
Sequence 1492	D61391	GENBANK	Sequence 1544	L38608	GENBANK
Sequence 1493	D63878	GENBANK	Sequence 1545	L38951	GENBANK
Sequence 1494	D63997	GENBANK	Sequence 1546	L42110	GENBANK
Sequence 1495	D82345	GENBANK	Sequence 1547	L47647	GENBANK
Sequence 1496	D83077	GENBANK	Sequence 1548	L49399	GENBANK
Sequence 1497	D86326	GENBANK	Sequence 1549	L77701	GENBANK
Sequence 1498	D86971	GENBANK	Sequence 1550	M10036	GENBANK
Sequence 1499	D86974	GENBANK	Sequence 1551	M10119	GENBANK
Sequence 1500	D86985	GENBANK	Sequence 1552	M11119	GENBANK
Sequence 1501	D87078	GENBANK	Sequence 1553	M13231	GENBANK
Sequence 1502	D87666	GENBANK	Sequence 1554	M13692	<b>GENBANK</b>
Sequence 1503	D87667	GENBANK	Sequence 1555	M14200	GENBANK
Sequence 1504	D88674	GENBANK	Sequence 1556	M15885	GENBANK
Sequence 1505	D89053	GENBANK	Sequence 1557	M15887	GENBANK

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Sequence 155		GENBA <b>NK</b>	Sequence 1610	U12778	GENBANK
Sequence 155		GENBANK	Sequence 1611	U14971	GENBANK
Sequence 156		GENBANK	Sequence 1612	U17714	GENBANK
Sequence 156		GENBA <b>NK</b>	Sequence 1613	U18914	GENBANK
Sequence 156	2 M20030	GENBANK	Sequence 1614	U21858	GENBANK
Sequence 156	3 M20372	GENBANK	Sequence 1615	U24169	GENBANK
Sequence 156	4 M21154	GENBANK	Sequence 1616	U25789	GENBANK
Sequence 156	5 M21895	<b>GENBANK</b>	Sequence 1617	U28249	GENBANK
Sequence 156	6 M21896	GENBANK	Sequence 1618	U30826	GENBANK
Sequence 156	7 M22382	GENBANK"	Sequence 1619	U31905	GENBANK
Sequence 156	8 M22538	GENBA <b>NK</b>	Sequence 1620	U33821	GENBANK
Sequence 156	9 M22590	GENBANK	Sequence 1621	U38654	GENBANK
Sequence 157		GENBANK	Sequence 1622	U38784	GENBANK
Sequence 157		GENBANK	Sequence 1623	U39360	GENBANK
Sequence 157		GENBANK	Sequence 1624	U41515	GENBANK
Sequence 157		GENBANK	Sequence 1625	U43188	GENBANK
Sequence 157		GENBANK	Sequence 1626	U48857	GENBANK
Sequence 157		GENBANK	Sequence 1627	U51004	GENBANK
Sequence 157		GENBANK	Sequence 1628	U51920	GENBANK
Sequence 157		GENBANK	Sequence 1629	U66469	GENBANK
Sequence 157		GENBANK	Sequence 1630	U66563	GENBANK
Sequence 157		GENBANK	Sequence 1631	U68140	GENBANK
Sequence 158		GENBANK	Sequence 1632	U70439	GENBANK
Sequence 158		GENBANK	Sequence 1633	U71363	GENBANK
Sequence 158		GENBANK	Sequence 1634	U77664	GENBANK
Sequence 158		GENBANK	Sequence 1635	U78516	GENBANK
Sequence 158		GENBANK	Sequence 1636	U792 <b>73</b>	GENBANK
Sequence 158		GENBANK	Sequence 1637	U80034	GENBANK
Sequence 158		GENBANK	Sequence 1638	U80456	GENBANK
Sequence 158		GENBANK	Sequence 1639	U84720	GENBANK
Sequence 158		GENBANK	Sequence 1640	U90 <b>551</b>	GENBANK
Sequence 1589	9 M64347	GENBANK	Sequence 1641	U97280	GENBANK
Sequence 1596		GENBANK	Sequence 1642	X00497	GENBANK
Sequence 159	1 M74777	GENBANK	Sequence 1643	X02544	GENBANK
Sequence 1592	2 M82882	GENBANK	Sequence 1644	X04707	GENBANK
Sequence 1593	3 M83822	GENBANK	Sequence 1645	X05332	GENBANK
Sequence 1594	M84443	GENBANK	Sequence 1646	X06747	GENBANK
Sequence 1598	M88163	<b>GENBANK</b>	Sequence 1647	X13238	GENBANK
Sequence 1596	M90104	GENBANK	Sequence 1648	X13839	GENBANK
Sequence 1597	7 M90309	GENBANK	Sequence 1649	X15183	GENBANK
Sequence 1598	M97856	<b>GENBANK</b>	Sequence 1650	X15187	GENBANK
Sequence 1599	S39329	<b>GENBANK</b>	Sequence 1651	X15722	GENBANK
Sequence 1600	S69189	GENBANK	Sequence 1652	X15729	GENBANK
Sequence 1601	S69272	GENBANK	Sequence 1653	X17620	GENBANK
Sequence 1602		GENBANK	Sequence 1654	X52943	GENBANK
Sequence 1603		GENBANK	Sequence 1655	X53280	GENBANK
Sequence 1604		GENBANK	Sequence 1656	X536 <b>05</b>	GENBANK
Sequence 1605		GENBANK	Sequence 1657	X56998	GENBANK
Sequence 1606		GENBANK	Sequence 1658	X59405	GENBANK
Sequence 1607		GENBANK	Sequence 1659	X60656	GENBANK
Sequence 1608		GENBANK	Sequence 1660	X60708	GE: BANK
Sequence 1609		GENBANK	Sequence 1661	X61100	GENBANK
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Sequence 1662	X66276	GENBANK	Sequence 1714	AA035001	dbEST
Sequence 1663	X69086	GENBANK	Sequence 1715	AA035090	dbEST
Sequence 1664	X73114	GENBANK	Sequence 1716	AA037138	dbEST
Sequence 1665	X74070	GENBANK	Sequence 1717	AA037216	dbEST
Sequence 1666	X74331	GENBANK	Sequence 1718	AA037828	dbEST
Sequence 1667	X75861	GENBANK	Sequence 1719	AA037877	dbEST
Sequence 1668	X76013	GENBANK	Sequence 1720	AA040037	dbEST
Sequence 1669	X76732	GENBANK	Sequence 1721	AA040820	dbEST
Sequence 1670	X81900	GENBANK	Sequence 1722	AA044591	dbEST
Sequence 1671	X82125	GENBANK	Sequence 1723	AA046808	dbEST
Sequence 1672	X82157	GENBANK	Sequence 1724	AA046973	dbEST
Sequence 1673	X90583	GENBANK	Sequence 1725	AA047729	dbEST
Sequence 1674	X94323	GENBANK	Sequence 1726	AA053522	dbEST
Sequence 1675	X95240	GENBANK	Sequence 1727	AA054120	dbEST
Sequence 1676	X95325	GENBANK	Sequence 1728	AA056034	dbEST
Sequence 1677	X97335	GENBANK	Sequence 1729	AA056482	dbEST
Sequence 1678	X98743	GENBANK	Sequence 1730	AA063501	dbEST
Sequence 1679	Y00052	GENBANK	Sequence 1731	AA069079	dbEST
Sequence 1680	Y00282	GENBANK	Sequence 1732	AA069693	dbEST
Sequence 1681	Y08915	GENBANK	Sequence 1733	AA069784	dbEST
Sequence 1682	Y08991	GENBANK	Sequence 1734	AA071084	dbEST
Sequence 1683	Y12860	GENBANK	Sequence 1735	AA074818	dbEST
Sequence 1684	Y13286	GENBANK	Sequence 1736	AA075165	dbEST
Sequence 1685	Z11227	GENBANK	Sequence 1737	AA075257	dbEST
Sequence 1686	Z26248	GENBANK	Sequence 1738	AA075530	dbEST
Sequence 1687	Z29067	GENBANK	Sequence 1739	AA077906	dbEST
Sequence 1688	Z31696	GENBANK	Sequence 1740	AA078904	dbEST
Sequence 1689	Z36819	GENBANK	Sequence 1741	AA080889	dbEST
Sequence 1690	Z48633	GENBANK	Sequence 1742	AA081751	dbEST
Sequence 1691	Z68747	GENBANK	Sequence 1743	AA082087	dbEST
Sequence 1692	Z69043	GENBANK	Sequence 1744	AA082754	dbEST
Sequence 1693	AA001319	dbEST	Sequence 1745	AA083096	dbEST
Sequence 1694	AA001792	dbEST	Sequence 1746	AA083410	dbEST
Sequence 1695	AA002181	dbEST	Sequence 1747	AA085459	dbEST
Sequence 1696	AA004609	dbEST	Sequence 1748	AA085632	dbEST
Sequence 1697	AA010480	dbEST	Sequence 1749	AA088187	dbEST
Sequence 1698	AA010828	dbEST	Sequence 1750	AA088372	dbEST
Sequence 1699	AA013090	dbEST	Sequence 1751	AA088755	dbEST
Sequence 1700	AA017469	dbEST	Sequence 1752	AA093974	dbEST
Sequence 1701	AA017500	dbEST	Sequence 1753	AA094656	dbEST
Sequence 1702	AA018347	dbEST	Sequence 1754	AA095478	dbEST
Sequence 1703	AA025373	dbEST	Sequence 1755	AA098876	dbEST
Sequence 1704	AA026983	dbEST	Sequence 1756	AA099080	dbEST
Sequence 1705	AA027862	dbEST	Sequence 1757	AA099602	dbEST
Sequence 1706	AA029889	dbEST	Sequence 1758	AA099976	dbEST
Sequence 1707	AA029947	dbEST	Sequence 1759	AA100707	dbEST
Sequence 1708	AA031293	dbEST	Sequence 1760	AA100885	dbEST
Sequence 1709	AA031762	dbEST	Sequence 1761	AA101237	dbEST
Sequence 1710	AA032221	dbEST	Sequence 1762	AA102605	dbEST
Sequence 1711	AA033869	dbEST	Sequence 1763	AA114945	dbEST
Sequence 1712	AA034186	dbEST	Sequence 1764	AA114952	dbEST
Sequence 1713	AA034516	dbEST	Sequence 1765	AA115021	dbEST
			•		

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Sequence 176	66 AA115492	2 dbEST	Sequence 1818	AA187809	dbEST
Sequence 176	57 AA115604	dbEST	Sequence 1819	AA188051	dbEST
Sequence 176	58 AA120852	dbEST	Sequence 1820	AA188140	dbEST
Sequence 176	59 AA122129	dbEST	Sequence 1821	AA190852	dbEST
Sequence 177	70 AA126 <mark>02</mark> 5	dbEST	Sequence 1822	AA190914	dbEST
Sequence 177	71 AA126 <mark>37</mark> 3	dbEST	Sequence 1823	AA191661	dbEST
Sequence 177	72 AA127336	dbEST	Sequence 1824	AA193247	dbEST
Sequence 177	73 AA127369	dbEST	Sequence 1825	AA195617	dbEST
Sequence 177	74 AA128114	dbEST	Sequence 1826	AA195831	dbEST
Sequence 17	75 AA130162	dbEST	Sequence 1827	AA196978	dbEST
Sequence 177	76 AA130229		Sequence 1828	AA203492	dbEST
Sequence 177			Sequence 1829	AA204854	dbEST
Sequence 177			Sequence 1830	AA205647	dbEST
Sequence 177			Sequence 1831	AA205870	dbEST
Sequence 178			Sequence 1832	AA206137	dbEST
Sequence 178			Sequence 1833	AA206214	dbEST
Sequence 178			Sequence 1834	AA207266	dbEST
Sequence 178			Sequence 1835	AA210741	dbEST
Sequence 178			Sequence 1836	AA210833	dbEST
Sequence 178			Sequence 1837	AA210840	dbEST
Sequence 178			Sequence 1838	AA218536	dbEST
Sequence 178			Sequence 1839	AA223774	dbEST
Sequence 178			Sequence 1840	AA224132	dbEST
Sequence 178			Sequence 1841	AA224751	dbEST
Sequence 179			Sequence 1842	AA224876	dbEST
Sequence 179			Sequence 1843	AA225002	
Sequence 179			Sequence 1844	AA225458	dbEST dbEST
Sequence 179			•	AA225857	
Sequence 179			Sequence 1845		dbEST
Sequence 179			Sequence 1846	AA226171 AA226275	dbEST
Sequence 179			Sequence 1847	AA226321	dbEST
Sequence 179			Sequence 1848		dbEST
Sequence 179			Sequence 1849 Sequence 1850	AA226611 AA226709	dbEST
Sequence 179			•	AA227873	dbEST
			Sequence 1851		dbEST
Sequence 180 Sequence 180			Sequence 1852	AA228440	dbEST
Sequence 180			Sequence 1853	AA229199	dbEST
•			Sequence 1854	AA229424	dbEST
Sequence 180			Sequence 1855	AA229611	dbEST
Sequence 180			Sequence 1856	AA230169	dbEST
Sequence 180			Sequence 1857	AA232734	dbEST
Sequence 180			Sequence 1858	AA232843	dbEST
Sequence 180			Sequence 1859	AA233428	dbEST
Sequence 180			Sequence 1860	AA234464	dbEST
Sequence 180			Sequence 1861	AA234818	dbEST
Sequence 181			Sequence 1862	AA235370	dbEST
Sequence 181			Sequence 1863	AA235802	dbEST
Sequence 181			Sequence 1864	AA236667	dbEST
Sequence 181			Sequence 1865	AA236904	dbEST
Sequence 181			Sequence 1866	AA242864	dbEST
Sequence 181			Sequence 1867	AA243230	dbEST
Sequence 181			Sequence 1868	AA243980	dbEST
Sequence 181	17 AA187288	dbEST	Sequence 1869	AA244052	dbEST

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Sequence 187		dbEST	Sequence 1922	AA313170	dbEST
Sequence 187		dbEST	Sequence 1923	AA3135 <b>34</b>	dbEST
Sequence 187	72 AA253 <b>384</b>	dbEST	Sequence 1924	AA3137 <b>79</b>	dbEST
Sequence 187	73 AA256 <b>591</b>	dbEST	Sequence 1925	AA314350	dbEST
Sequence 187	74 AA2583 <b>57</b>	dbEST	Sequence 1926	AA314599	dbEST
Sequence 187	75 AA2590 <b>62</b>	dbEST	Sequence 1927	AA314673	dbEST
Sequence 187		dbEST	Sequence 1928	AA315049	dbEST
Sequence 187	77 AA26 <b>2345</b>	dbEST	Sequence 1929	AA315189	dbEST
Sequence 187	78 AA278956	dbEST	Sequence 1930	AA315 <b>311</b>	dbEST
Sequence 187	'9 AA28084 <b>8</b>	dbEST	Sequence 1931	AA316 <b>181</b>	dbEST
Sequence 188	30 AA2839 <b>04</b>	dbEST	Sequence 1932	AA316632	dbEST
Sequence 188	31 AA284 <b>835</b>	dbEST	Sequence 1933	AA3179 <b>52</b>	dbEST
Sequence 188	32 AA284 <b>971</b>	dbEST	Sequence 1934	AA318100	dbEST
Sequence 188	3 AA285040	dbEST	Sequence 1935	AA320611	dbEST
Sequence 188	34 AA285 <b>290</b>	dbEST	Sequence 1936	AA322728	dbEST
Sequence 188	5 AA290591	dbEST	Sequence 1937	AA322 <b>922</b>	dbEST
Sequence 188	6 AA291680	dbEST	Sequence 1938	AA324683	dbEST
Sequence 188	7 AA292131	dbEST	Sequence 1939	AA325108	dbEST
Sequence 188	8 AA292866	dbEST	Sequence 1940	AA325220	dbEST
Sequence 188	9 AA293802	dbEST	Sequence 1941	AA329110	dbEST
Sequence 189	0 AA296465	dbEST	Sequence 1942	AA331359	dbEST
Sequence 189	1 AA297215	dbEST	Sequence 1943	AA332139	dbEST
Sequence 189	2 AA297512	dbEST	Sequence 1944	AA332551	dbEST
Sequence 189		dbEST	Sequence 1945	AA336798	dbEST
Sequence 189	4 AA298786	dbEST	Sequence 1946	AA340632	dbEST
Sequence 189		dbEST	Sequence 1947	AA342231	dbEST
Sequence 189	6 AA2995 <b>94</b>	dbEST	Sequence 1948	AA344376	dbEST
Sequence 189		dbEST	Sequence 1949	AA3459 <b>04</b>	dbEST
Sequence 189		dbEST	Sequence 1950	AA347 <b>340</b>	dbEST
Sequence 189		dbEST	Sequence 1951	AA351799	dbEST
Sequence 190		dbEST	Sequence 1952	AA354 <b>915</b>	dbEST
Sequence 190		dbEST	Sequence 1953	AA355 <b>119</b>	dbEST
Sequence 190		dbEST	Sequence 1954	AA357 <b>593</b>	dbEST
Sequence 190		dbEST	Sequence 1955	AA358169	dbEST
Sequence 190		dbEST	Sequence 1956	<b>AA</b> 3601 <b>90</b>	dbEST
Sequence 190		dbEST	Sequence 1957	<b>A</b> A36 <b>1997</b>	dbEST
Sequence 190		dbEST	Sequence 1958	AA3627 <b>31</b>	dbEST
Sequence 190		dbEST	Sequence 1959	<b>AA</b> 365 <b>229</b>	dbEST
Sequence 190		dbEST	Sequence 1960	AA3662 <b>69</b>	dbEST
Sequence 190		dbEST	Sequence 1961	AA369 <b>521</b>	dbEST
Sequence 191		dbEST	Sequence 1962	AA370110	dbEST
Sequence 191		dbEST	Sequence 1963	AA370122	dbEST
Sequence 191:		dbEST	Sequence 1964	AA3703 <b>37</b>	dbEST
Sequence 1913		dbEST	Sequence 1965	<b>A</b> A3730 <b>79</b>	dbEST
Sequence 1914		dbEST	Sequence 1966	AA374718	dbEST
Sequence 191		dbEST	Sequence 1967	AA375298	dbEST
Sequence 1916		dbEST	Sequence 1968	AA375312	dbEST
Sequence 1917		dbEST	Sequence 1969	AA3762 <b>92</b>	dbEST
Sequence 1918		dbEST	Sequence 1970	AA382481	dbEST
Sequence 1919		dbEST	Sequence 1971	AA384322	dbEST
Sequence 1920		dbEST	Sequence 1972	AA384529	dbEST
Sequence 1921	1 AA3126 <b>52</b>	dbEST	Sequence 1973	AA3847 <b>56</b>	dbEST

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Sequence 197		dbEST	Sequence 2026	AA451633	dbEST
Sequence 197		dbEST	Sequence 2027	AA453310	dbEST
Sequence 1976		dbEST	Sequence 2028	AA453562	dbEST
Sequence 197		dbEST	Sequence 2029	AA453924	dbEST
Sequence 1978		dbEST	Sequence 2030	AA455235	dbEST
Sequence 1979		dbEST	Sequence 2031	AA456344	dbEST
Sequence 1986	0 AA398619	dbEST	Sequence 2032	AA458516	dbEST
Sequence 198		dbEST	Sequence 2033	AA459321	dbEST
Sequence 1982	2 AA402000	dbEST	Sequence 2034	AA459517	dbEST
Sequence 1983	3 AA402238	dbEST	Sequence 2035	AA461128	dbEST
Sequence 1984		dbEST	Sequence 2036	AA464133	dbEST
Sequence 1985	5 AA406168	dbEST	Sequence 2037	AA464338	dbEST
Sequence 1986	6 AA406191	dbEST	Sequence 2038	AA464568	dbEST
Sequence 1987	7 AA418628	dbEST	Sequence 2039	AA468575	dbEST
Sequence 1988	B AA418685	dbEST	Sequence 2040	AA468839	dbEST
Sequence 1989	9 AA418824	dbEST	Sequence 2041	AA469199	dbEST
Sequence 1990	AA419011	dbEST	Sequence 2042	AA469406	dbEST
Sequence 1991	1 AA420625	dbEST	Sequence 2043	AA476263	dbEST
Sequence 1992	2 AA420645	dbEST	Sequence 2044	AA477173	dbEST
Sequence 1993		dbEST	Sequence 2045	AA477698	dbEST
Sequence 1994		dbEST	Sequence 2046	AA479614	dbEST
Sequence 1998	5 AA423957	dbEST	Sequence 2047	AA480420	dbEST
Sequence 1996		dbEST	Sequence 2048	AA480505	dbEST
Sequence 1997		dbEST	Sequence 2049	AA481355	dbEST
Sequence 1998		dbEST	Sequence 2050	AA482853	dbEST
Sequence 1999		dbEST	Sequence 2051	AA483565	dbEST
Sequence 2000		dbEST	Sequence 2052	AA485467	dbEST
Sequence 2001		dbEST	Sequence 2053	AA485767	dbEST
Sequence 2002		dbEST	Sequence 2054	AA487503	dbEST
Sequence 2003		dbEST	Sequence 2055	AA488280	dbEST
Sequence 2004		dbEST	Sequence 2056	AA489032	dbEST
Sequence 2005		dbEST	Sequence 2057	AA489410	dbEST
Sequence 2006		dbEST	Sequence 2058	AA489921	dbEST
Sequence 2007		dbEST	Sequence 2059	AA491081	dbEST
Sequence 2008		dbEST	Sequence 2060	AA491866	dbEST
Sequence 2009		dbEST	Sequence 2061	AA492026	dbEST
Sequence 2010		dbEST	Sequence 2062	AA492143	dbEST
Sequence 2011		dbEST	Sequence 2063	AA492280	dbEST
Sequence 2012		dbEST	Sequence 2064	AA493512	dbEST
Sequence 2013		dbEST	Sequence 2065	AA493742	dbEST
Sequence 2014		dbEST	Sequence 2066	AA493962	dbEST
Sequence 2015		dbEST	Sequence 2067	AA494344	dbEST
Sequence 2016		dbEST	Sequence 2068	AA501752	dbEST
Sequence 2017		dbEST	Sequence 2069	AA501945	dbEST
Sequence 2018		dbEST	Sequence 2070	AA502320	dbEST
Sequence 2019		dbEST	Sequence 2071	AA502979	dbEST
Sequence 2020		dbEST	Sequence 2072	AA503115	dbEST
Sequence 2021		dbEST	Sequence 2073	AA503224	dbEST
Sequence 2022		dbEST	Sequence 2074	AA503330	dbEST
Sequence 2023		dbEST	Sequence 2075	AA503611	dbEST
Sequence 2024		dbEST	Sequence 2076	AA503011	dbEST
Sequence 2025				AA504154 AA504265	
ocquence 2025	/ // <del>/44</del> 5003	dbEST	Sequence 2077	AA304203	dbEST

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Sequence 2078	AA504343	dbEST	Sequence 2130	AA564119	dbEST
Sequence 2079	AA505124	dbEST	Sequence 2131	AA565420	dbEST
Sequence 2080	AA505625	dbEST	Sequence 2132	AA568138	dbEST
Sequence 2081	AA506099	dbEST	Sequence 2133	AA568990	dbEST
Sequence 2082	AA507 <b>505</b>	dbEST	Sequence 2134	AA569765	dbEST
Sequence 2083	AA508804	dbEST	Sequence 2135	AA570519	dbEST
Sequence 2084	AA513597	dbEST	Sequence 2136	AA570665	dbEST
Sequence 2085	AA513940	dbEST	Sequence 2137	AA573975	dbEST
Sequence 2086	AA514490	dbEST	Sequence 2138	AA574223	dbEST
Sequence 2087	AA514775	dbEST	Sequence 2139	AA574397	dbEST
Sequence 2088	AA514991	dbEST	Sequence 2140	AA575933	dbEST
Sequence 2089	AA516531	dbEST	Sequence 2141	AA575935	dbEST
Sequence 2090	AA521142	dbEST	Sequence 2142	AA577800	dbEST
Sequence 2091	AA522849	dbEST	Sequence 2143	AA578577	dbEST
Sequence 2092	AA522850	dbEST	Sequence 2144	AA578773	dbEST
Sequence 2093	AA523252	dbEST	Sequence 2145	AA578904	dbEST
Sequence 2094	AA523498	dbEST	Sequence 2146	AA579617	dbEST
Sequence 2095	AA523751	dbEST	Sequence 2147	AA580279	dbEST
Sequence 2096	AA524064	dbEST	Sequence 2148	AA580399	dbEST
Sequence 2097	AA524258	dbEST	Sequence 2149	AA583449	dbEST
Sequence 2098	AA524285	dbEST	Sequence 2150	AA585110	dbEST
Sequence 2099	AA525269	dbEST	Sequence 2151	AA587468	dbEST
Sequence 2100	AA526403	dbEST	Sequence 2152	AA588848	dbEST
Sequence 2101	AA526430	dbEST	Sequence 2153	AA592908	dbEST
Sequence 2102	AA526834	dbEST	Sequence 2154	AA593385	dbEST
Sequence 2103	AA526869	dbEST	Sequence 2155	AA599669	dbEST
Sequence 2104	AA527342	dbEST	Sequence 2156	AA602961	dbEST
Sequence 2105	AA527730	dbEST	Sequence 2157	AA604619	dbEST
Sequence 2106	AA527805	dbEST	Sequence 2158	AA609394	dbEST
Sequence 2107	AA531068	dbEST	Sequence 2159	AA609641	dbEST
Sequence 2108	AA531487	dbEST	Sequence 2160	AA610581	dbEST
Sequence 2109	AA531563	dbEST	Sequence 2161	AA612573	dbEST
Sequence 2110	AA532398	dbEST	Sequence 2162	AA612864	dbEST
Sequence 2111	AA532934	dbEST	Sequence 2163	AA613891	dbEST
Sequence 2112	AA533 <b>57</b> 5	dbEST	Sequence 2164	AA614350	dbEST
Sequence 2113	AA533963	dbEST	Sequence 2165	AA620396	dbEST
Sequence 2114	AA534171	dbEST	Sequence 2166	AA620750	dbEST
Sequence 2115	AA534173	dbEST ·	Sequence 2167	AA622765	dbEST
Sequence 2116	AA534298	dbEST	Sequence 2168	AA627855	dbEST
Sequence 2117	AA534395	dbEST	Sequence 2169	AA630303	dbEST
Sequence 2118	AA536129	dbEST	Sequence 2170	AA631303	dbEST
Sequence 2119	AA541515	dbEST	Sequence 2171	AA631419	dbEST
Sequence 2120	AA541677	dbEST	Sequence 2172	AA631460	dbEST
Sequence 2121	AA553726	dbEST	Sequence 2173	AA631868	dbEST
Sequence 2122	AA553826	dbEST	Sequence 2174	AA634772	dbEST
Sequence 2123	AA554326	dbEST	Sequence 2175	AA635024	dbEST
Sequence 2124	AA557174	dbEST	Sequence 2176	AA639902	dbEST
Sequence 2125	AA557683	dbEST	Sequence 2177	AA640038	dbEST
Sequence 2126	AA558280	dbEST	Sequence 2178	AA640474	dbEST
Sequence 2127	AA558547	dbEST	Sequence 2179	AA641195	dbEST
Sequence 2128	AA558755	dbEST	Sequence 2180	AA644237	dbEST
Sequence 2129	AA559 <b>055</b>	dbEST	Sequence 2181	AA648318	dbEST
•			•		

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Sequence 2182	AA648757	dbEST	Sequence 2234	AA776740	dbEST
Sequence 2183	AA649012	dbEST	Sequence 2235	AA779728	dbEST
Sequence 2184	AA650114	dbEST	Sequence 2236	AA788779	dbEST
Sequence 2185	AA652167	dbEST	Sequence 2237	AA788875	dbEST
Sequence 2186	AA652478	dbEST	Sequence 2238	AA804544	dbEST
Sequence 2187	AA653601	dbEST	Sequence 2239	AA805228	dbEST
Sequence 2188	AA654875	dbEST	Sequence 2240	AA808492	dbEST
Sequence 2189	AA657411	dbEST	Sequence 2241	AA815305	dbEST
Sequence 2190	AA657488	dbEST	Sequence 2242	AA824401	dbEST
Sequence 2191	AA659388	dbEST	Sequence 2243	AA825768	dbEST
Sequence 2192	AA659719	dbEST	Sequence 2244	AA825940	dbEST
Sequence 2193	AA664480	dbEST	Sequence 2245	AA827875	dbEST
Sequence 2194	AA665209	dbEST	Sequence 2246	AA827878	dbEST
Sequence 2195	AA677531	dbEST	Sequence 2247	AA829236	dbEST
Sequence 2196	AA679791	dbEST	Sequence 2248	AA829547	dbEST
Sequence 2197	AA679935	dbEST	Sequence 2249	AA834540	dbEST
Sequence 2198	AA682244	dbEST	Sequence 2250	AA838003	dbEST
Sequence 2199	AA682252	dbEST	Sequence 2251	AA838410	dbEST
Sequence 2200	AA682283	dbEST	Sequence 2252	AA838681	dbEST
Sequence 2201	AA687216	dbEST	Sequence 2253	AA843326	dbEST
Sequence 2202	AA687336	dbEST	Sequence 2254	AA843592	dbEST
Sequence 2203	AA700232	dbEST	Sequence 2255	AA843706	dbEST
Sequence 2204	AA700876	dbEST	Sequence 2256	AA844070	dbEST
Sequence 2205	AA702888	dbEST	Sequence 2257	AA847656	dbEST
Sequence 2206	AA703396	dbEST	Sequence 2258	AA848129	dbEST
Sequence 2207	AA705433	dbEST	Sequence 2259	AA853183	dbEST
Sequence 2208	AA706764	dbEST	Sequence 2260	AA856719	dbEST
Sequence 2209	AA707607	dbEST	Sequence 2261	AA857476	dbEST
Sequence 2210	AA720578	dbEST	Sequence 2262	AA857936	dbEST
Sequence 2211	AA721375	dbEST	Sequence 2263	AA872606	dbEST
Sequence 2212	AA723525	dbEST	Sequence 2264	AA873182	dbEST
Sequence 2213	AA725400	dbEST	Sequence 2265	AA876668	dbEST
Sequence 2214	AA730534	dbEST	Sequence 2266	AA876897	dbEST
Sequence 2215	AA730780	dbEST	Sequence 2267	AA877789	dbEST
Sequence 2216	AA731556	dbEST	Sequence 2268	AA878212	dbEST
Sequence 2217	AA731700	dbEST	Sequence 2269	AA883580	dbEST
Sequence 2218	AA741297	dbEST	Sequence 2270	AA883901	dbEST
Sequence 2219	AA743975	dbEST	Sequence 2271	AA884098	dbEST
Sequence 2220	AA745694	dbEST	Sequence 2272	AA886818	dbEST
Sequence 2221	AA749045	dbEST	Sequence 2273	AA888589	dbEST
Sequence 2222	AA758348	dbEST	Sequence 2274	AA888665	dbEST
Sequence 2223	AA760738	dbEST	Sequence 2275	AA902388	dbEST
Sequence 2224	AA760894	dbEST	Sequence 2276	AA907575	dbEST
Sequence 2225	AA767154	dbEST	Sequence 2277	AA912796	dbEST
Sequence 2226	AA767457	dbEST	Sequence 2278	AA916635	dbEST
Sequence 2227	AA767764	dbEST	Sequence 2279	AA916753	dbEST
Sequence 2228	AA768348	dbEST	Sequence 2280	AA918023	dbEST
Sequence 2229	AA769434	dbEST	Sequence 2281	AA922237	dbEST
Sequence 2230	AA769( )6	dbEST	Sequence 2282	AA923278	dbEST
Sequence 2231	AA769722	dbEST	Sequence 2283	AA926992	dbEST
Sequence 2232	AA774869	dbEST	Sequence 2284	AA932252	dbEST
Sequence 2233	AA775918	dbEST	Sequence 2285	AA934368	dbEST

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Sequence 2286	AA934755	dbEST	Sequence 2338	AI087145	dbEST
Sequence 2287	AA935526	dbEST	Sequence 2339	AI087888	dbEST
Sequence 2288	AA935811	dbEST	Sequence 2340	A1088778	dbEST
Sequence 2289	AA939199	dbEST	Sequence 2341	A1090809	dbEST
Sequence 2290	AA939266	dbEST	Sequence 2342	AI092479	dbEST
Sequence 2291	AA948320	dbEST	Sequence 2343	A1094741	dbEST
Sequence 2292	AA954674	dbEST	Sequence 2344	A1097410	dbEST
Sequence 2293	AA962300	dbEST	Sequence 2345	AI114602	dbEST
Sequence 2294	AA968621	dbEST	Sequence 2346	AI114651	dbEST
Sequence 2295	AA968756	dbEST	Sequence 2347	Al126233	dbEST
Sequence 2296	AA969459	dbEST	Sequence 2348	AI126867	dbEST
Sequence 2297	AA969512	dbEST	Sequence 2349	AI128828	dbEST
Sequence 2298	AA973560	dbEST	Sequence 2350	Al130796	dbEST
Sequence 2299	AA977197	dbEST	Sequence 2351	Al133466	dbEST
Sequence 2300	AA977226	dbEST	Sequence 2352	AI133690	dbEST
Sequence 2301	AA978214	dbEST	Sequence 2353	AI138633	dbEST
Sequence 2302	AA983560	dbEST	Sequence 2354	AI138640	dbEST
Sequence 2303	AA987290	dbEST	Sequence 2355	AI140096	dbEST
Sequence 2304	AA991398	dbEST	Sequence 2356	AI141770	dbEST
Sequence 2305	AA995066	dbEST	Sequence 2357	AI142083	dbEST
Sequence 2306	AA995747	dbEST	Sequence 2358	AI143775	dbEST
Sequence 2307	AA996010	dbEST	Sequence 2359	AI149592	dbEST
Sequence 2308	AF034176	dbEST	Sequence 2360	AI151140	dbEST
Sequence 2309	AF063521	dbEST	Sequence 2361	Al161068	dbEST
Sequence 2310	AI015790	dbEST	Sequence 2362	AI174957	dbEST
Sequence 2311	Al018625	dbEST	Sequence 2363	AI174991	dbEST
Sequence 2312	Al022687	dbEST	Sequence 2364	AI183358	dbEST
Sequence 2313	A1024549	dbEST	Sequence 2365	AI183447	dbEST
Sequence 2314	Al025419	dbEST	Sequence 2366	AI184562	dbEST
Sequence 2315	AI026767	dbEST	Sequence 2367	AI191244	dbEST
Sequence 2316	AI027536	dbEST	Sequence 2368	AI192767	dbEST
Sequence 2317	AI032392	dbEST	Sequence 2369	AI198311	dbEST
Sequence 2318	AI032838	dbEST	Sequence 2370	Al198986	dbEST
Sequence 2319	AI038692	dbEST	Sequence 2371	Al199094	dbEST
Sequence 2320	AI040598	dbEST	Sequence 2372	AI200038	dbEST
Sequence 2321	AI050871	dbEST	Sequence 2373	Al200110	dbEST
Sequence 2322	AI051210	dbEST	Sequence 2374	AI204090	dbEST
Sequence 2323	AI052333	dbEST	Sequence 2375	AI206199	dbEST
Sequence 2324	AI055939	dbEST	Sequence 2376	AI206344	dbEST
Sequence 2325	AI061649	dbEST	Sequence 2377	AI207367	dbEST
Sequence 2326	AI064691	dbEST	Sequence 2378	AI208947	dbEST
Sequence 2327	A1064844	dbEST	Sequence 2379	AI215841	dbEST
Sequence 2328	AI065092	dbEST	Sequence 2380	AI216969	dbEST
Sequence 2329	Al073465	dbEST	Sequence 2381	AI216988	dbEST
Sequence 2330	AI075324	dbEST	Sequence 2382	AI217003	dbEST
Sequence 2331	AI079558	dbEST	Sequence 2383	AI217012	dbEST
Sequence 2332	AI080267	dbEST	Sequence 2384	AI217021	dbEST
Sequence 2333	AI080480	dbEST	Sequence 2385	AI217023	dbEST
Sequence 2334	AI082168	dbEST	Sequence 2386	AI217425	dbEST
Sequence 2335	AI083814	dbEST	Sequence 2387	AI218376	dbEST
Sequence 2336	A1084604	dbEST	Sequence 2388	AI218882	dbEST
Sequence 2337	Al085165	dbEST	Sequen <b>ce</b> 2389	Al219324	dbEST

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Sequence 2390	Al221829	dbEST	Sequence 2442	Al347506	dbEST
Sequence 2391	AI223085	dbEST	Sequence 2443	Al356718	dbEST
Sequence 2392	Al224519	dbEST	Sequence 2444	Al362403	dbEST
Sequence 2393	Al225043	dbEST	Sequence 2445	Al362462	dbEST
Sequence 2394	Al239822	dbEST	Sequence 2446	Al364053	dbEST
Sequence 2395	Al240872	dbEST	Sequence 2447	Al367350	dbEST
Sequence 2396	A1243423	dbEST	Sequence 2448	Al369174	dbEST
Sequence 2397	Al243780	dbEST	Sequence 2449	Al376513	dbEST
Sequence 2398	Al248277	dbEST	Sequence 2450	Al376865	dbEST
Sequence 2399	Al248736	dbEST	Sequence 2451	Al377194	dbEST
Sequence 2400	Al249000	dbEST	Sequence 2452	Al391631	dbEST
Sequence 2401	Al249797	dbEST	Sequence 2453	Al394646	dbEST
Sequence 2402	Al253330	dbEST	Sequence 2454	Al400752	dbEST
Sequence 2403	Al253335	dbEST	Sequence 2455	Al401840	dbEST
Sequence 2404	Al253338	dbEST	Sequence 2456	Al420227	dbEST
Sequence 2405	Al253379	dbEST	Sequence 2457	Al431814	dbEST
Sequence 2406	Al253436	dbEST	Sequence 2458	Al433818	dbEST
Sequence 2407	Al262486	dbEST	Sequence 2459	Al439774	dbEST
Sequence 2408	Al265776	dbEST	Sequence 2460	Al452703	dbEST
Sequence 2409	Al267162	dbEST	Sequence 2461	Al469038	dbEST
Sequence 2410	Al267185	dbEST	Sequence 2462	Al472525	dbEST
Sequence 2411	Al267276	dbEST	Sequence 2463	Al474466	dbEST
Sequence 2412	Al267282	dbEST	Sequence 2464	Al476318	dbEST
Sequence 2413	Al267307	dbEST	Sequence 2465	Al478365	dbEST
Sequence 2414	Al267454	dbEST	Sequence 2466	Al479439	dbEST
Sequence 2415	Al267502	dbEST	Sequence 2467	Al494299	db <b>EST</b>
Sequence 2416	AI268328	dbEST	Sequence 2468	Al499393	dbEST
Sequence 2417	Al268405	dbEST	Sequence 2469	Al499986	dbEST
Sequence 2418	Al268860	dbEST	Sequence 2470	Al521100	dbEST
Sequence 2419	Al269060	dbEST	Sequence 2471	AI523506	dbEST
Sequence 2420	Al270511	dbEST	Sequence 2472	AI525231	dbEST
Sequence 2421	Al272941	dbEST	Sequence 2473	AI525454	dbEST
Sequence 2422	AI275175	dbEST	Sequence 2474	AI536024	dbEST
Sequence 2423	Al279838	dbEST	Sequence 2475	AI539349	dbEST
Sequence 2424	Al282277	dbEST	Sequence 2476	AI547146	dbEST
Sequence 2425	Al285488	dbEST	Sequence 2477	AI554016	dbEST
Sequence 2426	Al289220	dbEST	Sequence 2478	Al557112	dbEST
Sequence 2427	Al291247	dbEST	Sequence 2479	AI557116	dbEST
Sequence 2428	Al299524	dbEST	Sequence 2480	AI557182	dbEST
Sequence 2429	Al302137	dbEST	Sequence 2481	AI557225	dbEST
Sequence 2430	Al302799	dbEST	Sequence 2482	AI557231	dbEST
Sequence 2431	Al304947	dbEST	Sequence 2483	AI557235	dbEST
Sequence 2432	AI307402	dbEST	Sequence 2484	AI557246	dbEST
Sequence 2433	AI307515	dbEST	Sequence 2485	AI557495	dbEST
Sequence 2434	Al318280	dbEST	Sequence 2486	AI557599	dbEST
Sequence 2435	Al334833	dbEST	Sequence 2487	AI557626	dbEST
Sequence 2436	Al334964	dbEST	Sequence 2488	AI560958	dbEST
Sequence 2437	Al335153	dbEST	Sequence 2489	AI561122	dbEST
Sequence 2438	Al335554	dbEST	Sequence 2490	AI564475	dbEST
Sequence 2439	Al336147	dbEST	Sequence 2491	AI564487	dbEST
Sequence 2440	AI338335	dbEST	Sequence 2492	AI580160	dbEST
Sequence 2441	Al345416	dbEST	Sequence 2493	AI580162	dbEST

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Sequence 2494	AI580927	db <b>EST</b>	Sequence 2546	Al7688 <b>54</b>	dbEST
Sequence 2495	A1589476	dbEST	Sequence 2547	AI768877	dbEST
Sequence 2496	AI591088	dbEST	Sequence 2548	A1770098	dbEST
Sequence 2497	A1620673	dbEST	Sequence 2549	Al791179	dbEST
Sequence 2498	Al621153	dbEST	Sequence 2550	Al791419	dbEST
Sequence 2499	AI623315	dbEST	Sequence 2551	A1796120	dbEST
Sequence 2500	Al623804	dbEST	Sequence 2552	Al797042	dbEST
Sequence 2501	AI625404	dbEST	Sequence 2553	A1800250	dbEST
Sequence 2502	A1625747	dbEST	Sequence 2554	AI800673	dbEST
Sequence 2503	AI627356	dbEST	Sequence 2555	AI804662	dbEST
Sequence 2504	AI627542	dbEST	Sequence 2556	AI805522	dbEST
Sequence 2505	Al630104	dbEST	Sequence 2557	AI809314	dbEST
Sequence 2506	Al630188	dbEST	Sequence 2558	AI810002	dbEST
Sequence 2507	AI630634	dbEST	Sequence 2559	AI815499	dbEST
Sequence 2508	AI632523	dbEST	Sequence 2560	AI816322	dbEST
Sequence 2509	A1632534	dbEST	Sequence 2561	AI821437	dbEST
Sequence 2510	Al636943	dbEST	Sequence 2562	AI826105	dbEST
Sequence 2511	Al638414	dbEST	Sequence 2563	AI830758	dbEST
Sequence 2512	Al650721	dbEST	Sequence 2564	AI831462	dbEST
Sequence 2513	Al651195	dbEST	Sequence 2565	AI863042	dbEST
Sequence 2514	Al653319	dbEST	Sequence 2566	A1864870	dbEST
Sequence 2515	Al653621	dbEST	Sequence 2567	AI867463	dbEST
Sequence 2516	Al659898	dbEST	Sequence 2568	Al873501	dbEST
Sequence 2517	Al669210	dbEST	Sequence 2569	Al878926	dbEST
Sequence 2518	AI670009	dbEST	Sequence 2570	Al879040	dbEST
Sequence 2519	A1680343	dbEST	Sequence 2571	A1879248	dbEST
Sequence 2520	Al683431	dbEST	Sequence 2572	Al8849 <b>63</b>	dbEST
Sequence 2521	Al688098	dbEST	Sequence 2573	A1887875	dbEST
Sequence 2522	Al689722	dbEST	Sequence 2574	Al914133	dbEST
Sequence 2523	Al689762	dbEST	Sequence 2575	Al923011	dbEST
Sequence 2524	Al690157	dbEST	Sequence 2576	Al923978	dbEST
Sequence 2525	Al693561	dbEST	Sequence 2577	A1929113	dbEST
Sequence 2526	AI700642	dbEST	Sequence 2578	AI936548	dbEST
Sequence 2527	A1709356	dbEST	Sequence 2579	AI936 <b>562</b>	dbEST
Sequence 2528	A1732680	dbEST	Sequence 2580	AI948503	dbEST
Sequence 2529	A1734209	dbEST	Sequence 2581	A1949422	dbEST
Sequence 2530	A1738605	dbEST	Sequence 2582	Al951118	dbEST
Sequence 2531	A1740900	dbEST	Sequence 2583	Al951970	dbEST
Sequence 2532	Al741486	dbEST	Sequence 2584	Al953711	dbEST
Sequence 2533	AJ741633	dbEST	Sequence 2585	Al969567	dbEST
Sequence 2534	Al742115	dbEST	Sequence 2586	AI972279	dbEST
Sequence 2535	AJ743303	dbEST	Sequence 2587	A1972479	dbEST
Sequence 2536	AI743852	dbEST	Sequence 2588	Al979288	dbEST
Sequence 2537	Al750682	dbEST	Sequence 2589	Al986240	dbEST
Sequence 2538	Al753280	dbEST	Sequence 2590	AL036499	dbEST
Sequence 2539	Al753951	dbEST	Sequence 2591	AL037031	dbEST
Sequence 2540	AI755085	dbEST	Sequence 2592	AL037226	dbEST
Sequence 2541	Al761291	dbEST	Sequence 2593	AL037707	dbEST
Sequence 2542	AI761469	dbEST	Sequence 2594	AL037798	dbEST
Sequence 2543	A1761986	dbEST	Sequence 2595	AL037828	dbEST
Sequence 2544	AI764962	dbEST	Sequence 2596	AL037847	dbEST
Sequence 2545	AI768624	dbEST	Sequence 2597	AL038 <b>802</b>	dbEST
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Sequence 2598	AL038985	dbEST	Sequence 2650	C18232	dbEST
Sequence 2599	AL03 <b>9254</b>	dbEST	Sequence 2651	D50977	dbEST
Sequence 2600	AL039293	dbEST	Sequence 2652	D54335	dbEST
Sequence 2601	AL039542	dbEST	Sequence 2653	D58694	dbEST
Sequence 2602	AL040692	dbEST	Sequence 2654	D60970	dbEST
Sequence 2603	AL041050	dbEST	Sequence 2655	D81 <b>855</b>	dbEST
Sequence 2604	AL041176	dbEST	Sequence 2656	F02595	dbEST
Sequence 2605	AL042234	dbEST	Sequence 2657	F084 <b>93</b>	dbEST
Sequence 2606	AL042613	db <b>EST</b>	Sequence 2658	F27229	dbEST
Sequence 2607	AL043034	dbEST	Sequence 2659	H03650	dbEST
Sequence 2608	AL043506	dbEST	Sequence 2660	H03785	dbEST
Sequence 2609	AL043864	d <b>bEST</b>	Sequence 2661	H06621	dbEST
Sequence 2610	AL044891	dbEST	Sequence 2662	H06944	dbEST
Sequence 2611	AL045213	dbEST	Sequence 2663	H09332	dbEST
Sequence 2612	AL045331	dbEST	Sequence 2664	H12519	dbEST
Sequence 2613	AL045538	d <b>bEST</b>	Sequence 2665	H24176	dbEST
Sequence 2614	AL048890	dbEST	Sequence 2666	H24856	dbEST
Sequence 2615	AL118999	dbEST	Sequence 2667	H26492	dbEST
Sequence 2616	AL120381	dbEST	Sequence 2668	H40738	dbEST
Sequence 2617	AW002185	dbEST	Sequence 2669	H47252	dbEST
Sequence 2618	AW003247	dbEST	Sequence 2670	H47363	dbEST
Sequence 2619	AW008473	dbEST	Sequence 2671	H49329	dbEST
Sequence 2620	AW015055	dbEST	Sequence 2672	H49828	dbEST
Sequence 2621	AW020699	dbEST	Sequence 2673	H57382	dbEST
Sequence 2622	AW022599	dbEST	Sequence 2674	H57485	dbEST
Sequence 2623	AW027339	dbEST	Sequence 2675	F.60344	dbEST
Sequence 2624	AW028594	dbEST	Sequence 2676	H73387	dbEST
Sequence 2625	AW028606	dbEST	Sequence 2677	H82380	dbEST
Sequence 2626	AW044547	dbEST	Sequence 2678	H87420	dbEST
Sequence 2627	AW067913	dbEST	Sequence 2679	H93552	dbEST
Sequence 2628	AW071696	dbEST	Sequence 2680	H94179	dbEST
Sequence 2629	AW076087	dbEST	Sequence 2681	H95161	dbEST
Sequence 2630	AW078747	dbEST	Sequence 2682	N27899	dbEST
Sequence 2631	AW084017	dbEST	Sequence 2683	N32072	dbEST
Sequence 2632	AW102951	dbEST	Sequence 2684	N34186	dbEST
Sequence 2633	AW104561	dbEST	Sequence 2685	N44337	dbEST
Sequence 2634	AW118457	db <b>EST</b>	Sequence 2686	N45167	dbEST
Sequence 2635	AW118595	dbEST	Sequence 2687	N46947	dbEST
Sequence 2636	AW131770	dbEST	Sequence 2688	N52271	dbEST
Sequence 2637	AW134772	dbEST	Sequence 2689	N52432	dbEST
Sequence 2638	AW135359	dbEST	Sequence 2690	N53033	dbEST
Sequence 2639	AW138731	dbEST	Sequence 2691	N54973	dbEST
Sequence 2640	AW140056	dbEST	Sequence 2692	N64576	dbEST
Sequence 2641	AW148884	db <b>EST</b>	Sequence 2693	N72648	dbEST
Sequence 2642	AW150276	dbEST	Sequence 2694	N72922	dbEST
Sequence 2643	AW152311	dbEST	Sequence 2695	N74267	dbEST
Sequence 2644	AW157060	db <b>EST</b>	Sequence 2696	N76672	dbEST
Sequence 2645	AW157303	db <b>EST</b>	Sequence 2697	N79427	dbEST
Sequence 2646	AW160447	db <b>EST</b>	Sequence 2698	N92160	dbEST
Sequence 2647	AW167579	dbEST	Sequence 2699	N94098	dbEST
Sequence 2648	AW176926	dbEST	Sequence 2700	R10438	dbEST
Sequence 2649	C15322	db <b>EST</b>	Sequence 2701	R10553	dbEST

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Sequence 2702	R13850	dbEST	Sequence 2754	Z45274	ANECT
Sequence 2703	R24803	dbEST	Sequence 2755	Q44278	dbEST NUCPATENT
Sequence 2704	R28584	dbEST	Sequence 2756	Q46540	
Sequence 2705	R34443	dbEST	Sequence 2757	Q49943	NUCPATENT
Sequence 2706	R35797	dbEST	Sequence 2758	Q53478	NUCPATENT
Sequence 2707	R45144	dbEST	Sequence 2759	T18813	NUCPATENT
Sequence 2708	R48890	dbEST	Sequence 2760	T28638	NUCPATENT
Sequence 2709	R70639	dbEST	Sequence 2761	T43474	NUCPATENT
Sequence 2710	R76235	dbEST	Sequence 2762	T59274	NUCPATENT
Sequence 2711	R76244	dbEST	Sequence 2763	V26460	NUCPATENT
Sequence 2712	R77186	dbEST	Sequence 2764	V20400 V41451	NUCPATENT
Sequence 2713	R80279	dbEST	Sequence 2765		NUCPATENT
Sequence 2714	R80852	dbEST	Sequence 2766	V47539	NUCPATENT
Sequence 2715	R84308	dbEST	Sequence 2767	V49578	NUCPATENT
Sequence 2716	R86265	dbEST		V58523	NUCPATENT
Sequence 2717	R87877	dbEST	Sequence 2768	V58544	NUCPATENT
Sequence 2718	R89611	dbEST	Sequence 2769	V58628	NUCPATENT
Sequence 2719	R92548	dbEST	Sequence 2770	V58685	NUCPATENT
Sequence 2720	R95821	dbEST	Sequence 2771	V58761	NUCPATENT
Sequence 2721	T10145	dbEST	Sequence 2772	V59638	NUCPATENT
Sequence 2722	T18977		Sequence 2773	V61487	NUCPATENT
Sequence 2723	T32212	dbEST	Sequence 2774	V62427	NUCPATENT
Sequence 2724	T41232	dbEST	Sequence 2775	V62428	NUCPATENT
Sequence 2725	T52823	dbEST	Sequence 2776	V62430	NUCPATENT
Sequence 2726	T53424	dbEST	Sequence 2777	V68992	NUCPATENT
Sequence 2727		dbEST	Sequence 2778	V84497	NUCPATENT
Sequence 2728	T54095	dbEST	Sequence 2779	V84510	NUCPATENT
Sequence 2729	T57386	dbEST	Sequence 2780	V87984	NUCPATENT
Sequence 2730	T64630	dbEST	Sequence 2781	V90138	NUCPATENT
Sequence 2731	T65098	dbEST	Sequence 2782	X00630	NUCPATENT
_ `	T67265	dbEST	Sequence 2783	X00632	NUCPATENT
Sequence 2732 Sequence 2733	T78615	dbEST	Sequence 2784	X03841	NUCPATENT
	T83709	dbEST	Sequence 2785	X04407	NUCPATENT
Sequence 2734	T93613	dbEST	Sequence 2786	X25445	NUCPATENT
Sequence 2735	T99715	dbEST	Sequence 2787	X26850	NUCPATENT
Sequence 2736	U46309	dbEST	Sequence 2788	X27262	NUCPATENT
Sequence 2737	W04421	dbEST	Sequence 2789	X27278	NUCPATENT
Sequence 2738	W19519	dbEST	Sequence 2790	X37486	NUCPATENT
Sequence 2739	W25008	dbEST	Sequence 2791	X40477	NUCPATENT
Sequence 2740	W31992	dbEST	Sequence 2792	X40518	NUCPATENT
Sequence 2741	W39298	dbEST	Sequence 2793	X40711	NUCPATENT
Sequence 2742	W42653	dbEST	Sequence 2794	X41185	NUCPATENT
Sequence 2743	W52896	dbEST	Sequence 2795	X90444	NUCPATENT
Sequence 2744	W56077	dbEST	Sequence 2796	X97772	NUCPATENT
Sequence 2745	W63676	dbEST	Sequence 2797	X98891	NUCPATENT
Sequence 2746	W68028	dbEST	Sequence 2798	Z11491	NUCPATENT
Sequence 2747	W86160	dbEST	Sequence 2799	Z14468	NUCPATENT
Sequence 2748	<b>W8</b> 786 <b>5</b>	dbEST	Sequence 2800	Z15132	NUCPATENT
Sequence 2749	W88823	dbEST	Sequence 2801	Z16412	NUCPATENT
Sequence 2750	W90224	dbEST	Sequence 2802	AC28312	PREPATNUC
Sequence 2751	W91890	dbEST	Sequence 2803	AC31081	PREPATNUC
Sequence 2752	Z43146	dbEST	Sequence 2804	AC31093	PREPATNUC
Sequence 2753	Z43843	dbEST	Sequence 2805	AC33927	PREPATNUC
			•		

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Sequence #	Accession #	Database	Sequence 2857	AB019524	GENBANK
Sequence 2806	A06846	GENBANK	Sequence 2858	AB019564	GENBANK
Sequence 2807	A17546	GENBANK	Sequence 2859		GENBANK
Sequence 2808	A18657	GENBANK	Sequence 2860	AB019987	GENBANK
Sequence 2809	AB000095	GENBANK	Sequence 2861	AB020650	GENBANK
Sequence 2810	AB000468	GENBANK	Sequence 2862	AB020673	GENBANK
Sequence 2811	AB002303	GENBANK	Sequence 2863	AB020682	GENBANK
Sequence 2812	AB002311	GENBANK	Sequence 2864	AB020686	GENBANK
Sequence 2813	AB002319	GENBANK	Sequence 2865	AB020880	GENBANK
Sequence 2814	AB002330	GENBANK	Sequence 2866	AB021663	GENBANK
Sequence 2815	AB002331	GENBANK	Sequence 2867	AB023198	GENBANK
Sequence 2816	AB002365	GENBANK	Sequence 2868	AB023205	GENBANK
Sequence 2817	AB002366	GENBANK	Sequence 2869	AB023215	GENBANK
Sequence 2818	AB002370	GENBANK	Sequence 2870	AB024704	GENBANK
Sequence 2819	AB002380	GENBANK	Sequence 2871	AB027466	GENBANK
Sequence 2820	AB002386	GENBANK	Sequence 2872	AB028624	GENBANK
Sequence 2821	AB004788	GENBANK	Sequence 2873	AB028951	GENBANK
Sequence 2822	AB004851	GENBANK	Sequence 2874	AB028995	GENBANK
Sequence 2823	AB006625	GENBANK	Sequence 2875	AB029008	GENBANK
Sequence 2824	AB007891	GENBANK	Sequence 2876	AB029025	GENBANK
Sequence 2825	AB007899	GENBANK	Sequence 2877	AB029290	GENBANK
Sequence 2826	AB007916	GENBANK	Sequence 2878	AB032969	GENBANK
Sequence 2827	AB007929	GENBANK	Sequence 2879	AB032983	GENBANK
Sequence 2828	AB007932	GENBANK	Sequence 2880	AB032994	GENBANK
Sequence 2829	AB007935	GENBANK	Sequence 2881	AB033007	GENBANK
Sequence 2830	AB007956	GENBANK	Sequence 2882	AB033010	GENBANK
Sequence 2831	AB007957	GENBANK	Sequence 2883	AB033033	GENBANK
Sequence 2832	AB010427	GENBANK	Sequence 2884	AB033059	GENBANK
Sequence 2833	AB011087	GENBANK	Sequence 2885	AB033070	GENBANK
Sequence 2834	AB011103	GENBANK	Sequence 2886	AB033091	GENBANK
Sequence 2835	AB011108	GENBANK	Sequence 2887	AB033899	GENBANK
Sequence 2836	AB011147	GENBANK	Sequence 2888	AF000152	GENBANK
Sequence 2837	AB011164	GENBANK	Sequence 2889	AF000381	GENBANK
Sequence 2838	AB011169	GENBANK	Sequence 2890	AF001893	GENBANK
Sequence 2839	AB011170	GENBANK	Sequence 2891	AF002210	GENBANK
Sequence 2840	AB013918	GENBANK	Sequence 2892	AF002668	GENBANK
Sequence 2841	AB014511	GENBANK	Sequence 2893	AF003341	GENBANK
Sequence 2842	AB014533 -	GENBANK	Sequence 2894	AF004561	GENBANK
Sequence 2843	AB014536	GENBANK	Sequence 2895	AF004813	GENBANK
Sequence 2844	AB014542	GENBANK	Sequence 2896	AF005038	GENBANK
Sequence 2845	AB014577	GENBANK	Sequence 2897	AF005043	GENBANK
Sequence 2846	AB014888	GENBANK	Sequence 2898	AF005050	GENBANK
Sequence 2847	AB015907	GENBANK	Sequence 2899	AF006086	GENBANK
Sequence 2848	AB016533	GENBANK	Sequence 2900	AF006514	GENBANK
Sequence 2849	AB017430	GENBANK	Sequence 2901	AF007144	GENBANK
Sequence 2850	AB018274	GENBANK	Sequence 2902	AF007151	GENBANK
Sequence 2851	AB018320	GENBANK	Sequence 2903	AF007216	GENBANK
Sequence 2852	AB018324	GENBANK	Sequence 2904	AF009620	GENBANK
Sequence 2853	AB018330	GENBANK	Sequence 2905	AF010233	GENBANK
Sequence 2854	AB013334	GENBANK	Sequence 2906	AF010313	GENBANK
Sequence 2855	AB018338	GENBANK	Sequence 2907	AF011904	GENBANK
Sequence 2856	AB018353	GENBANK	Sequence 2908	AF012072	GENBANK

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Sequence 2909	9 AF013168	GENBANK	Sequ	ience 2961	AF070561	GENBANK
Sequence 291	O AF013759	GENBANK	Sequ	ience 2962	AF070650	
Sequence 291	1 AF013970	GENBANK	Sequ	ience 2963	AF070657	
Sequence 2912	2 AF015283	GENBANK		ence 2964	AF070665	GENBANK
Sequence 291:	3 AF020038	GENBANK	-	ience 2965	AF071202	GENBANK
Sequence 291	4 AF020202	GENBANK		ence 2966	AF072810	GENBANK
Sequence 291		GENBANK	•	ience 2967	AF073310	GENBANK
Sequence 2916		GENBANK		ence 2968	AF075587	GENBANK
Sequence 291		GENBANK		ience 2969	AF076929	GENBANK
Sequence 2918		GENBANK		ence 2970	AF077030	GENBANK
Sequence 2919		GENBANK	•	ence 2971	AF077043	GENBANK
Sequence 2920		GENBANK	•	ence 2972	AF077202	GENBANK
Sequence 292		GENBANK	·	ence 2973	AF077599	GENBANK
Sequence 2922		GENBANK		ence 2974	AF081192	GENBANK
Sequence 2923		GENBANK	•	ence 2975	AF081282	GENBANK
Sequence 2924		GENBANK	·	ence 2976	AF082868	GENBANK
Sequence 2925		GENBANK	•	ence 2977	AF083190	GENBANK
Sequence 2926		GENBANK	·	ence 2978	AF083395	GENBANK
Sequence 2927		GENBANK	•	ence 2979	AF085359	GENBANK
Sequence 2928		GENBANK	•	ence 2980	AF085362	GENBANK
Sequence 2929		GENBANK	•	ence 2981	AF085844	GENBANK
Sequence 2930		GENBANK		ence 2982	AF085893	GENBANK
Sequence 2931		GENBANK	•	ence 2983	AF085917	GENBANK
Sequence 2932		GENBANK	•	ence 2984	AF086183	GENBANK
Sequence 2933		GENBANK	· ·	ence 2985	AF087481	GENBANK
Sequence 2934		GENBANK	•	ence 2986	AF092038	GENBANK
Sequence 2935		GENBANK	•	ence 2987	AF092905	GENBANK
Sequence 2936		GENBANK	·	ence 2988	AF100741	GENBANK
Sequence 2937		GENBANK	•	ence 2989	AF100755	GENBANK
Sequence 2938		GENBANK	•	ence 2990	AF102265	GENBANK
Sequence 2939		GENBANK	-	ence 2991	AF102988	GENBANK
Sequence 2940		GENBANK	•	ence 2992	AF103801	GENBANK
Sequence 2941		GENBANK	•	ence 2993	AF103804	GENBANK
Sequence 2942		GENBANK	•	ence 2994	AF103907	GENBANK
Sequence 2943	3 AF054187	GENBANK	•	ence 2995	AF104013	GENBANK
Sequence 2944		GENBANK		ence 2996	AF110647	GENBANK
Sequence 2945		GENBANK	·	ence 2997	AF117949	GENBANK
Sequence 2946	AF055008	GENBANK		ence 2998	AF119121	GENBANK
Sequence 2947	7 AF055022	GENBANK	•	ence 2999	AF126424	GENBANK
Sequence 2948	3 AF055025	GENBANK	Sequ	ence 3000	AF126782	GENBANK
Sequence 2949		GENBANK		ence 3001	AF128406	GENBANK
Sequence 2950	AF056087	GENBANK		ence 3002	AF131774	GENBANK
Sequence 2951	AF057160	GENBANK	·	ence 3003	AF131856	GENBANK
Sequence 2952	2 AF058319	GENBANK	·	ence 3004	AF132000	GENBANK
Sequence 2953	AF060152	GENBANK		ence 3005	AF132937	GENBANK
Sequence 2954		GENBANK	•	ence 3006	AF133845	GENBANK
Sequence 2955		GENBANK	•	ence 3007	AF136450	GENBANK
Sequence 2956		GENBANK	•	ence 3008	AF139461	GENBANK
Sequence 2957		GENBANK	•	ence 3009	AF143235	GENBANK
Sequence 2958		GENBANK		ence 3010	AF144755	GENBANK
Sequence 2959		GENBANK		ence 3011	AF145029	GENBANK
Sequence 2960		GENBANK		ence 3012	AF145316	GENBANK
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Sequence 3013	AF147367	GENBANK	Sequence 30	55 AL080209	GENBANK
Sequence 3014	AF150087	GENBANK	Sequence 30		GENBANK
Sequence 3015	AF151103	GENBANK	Sequence 30		GENBANK
Sequence 3016	AF151840	GENBANK	Sequence 30		GENBANK
Sequence 3017	AF151861	GENBANK	Sequence 30		GENBANK
Sequence 3018	AF151870	GENBANK	Sequence 30		GENBANK
Sequence 3019	AF151902	GENBANK	Sequence 30		GENBANK
Sequence 3020	AF155100	GENBANK	Sequence 30		GENBANK
Sequence 3021	AF155113	GENBANK	Sequence 30		GENBANK
Sequence 3022	AF155330	GENBANK	Sequence 30		GENBANK
Sequence 3023	AF155568	GENBANK	Sequence 30		GENBANK
Sequence 3024	AF157028	GENBANK	Sequence 30		GENBANK
Sequence 3025	AF159056	GENBANK	Sequence 30		GENBANK
Sequence 3026	AF165281	GENBANK	Sequence 30		GENBANK
Sequence 3027	AF168956	GENBANK	Sequence 30		GENBANK
Sequence 3028	AF169481	GENBANK	Sequence 30		GENBANK
Sequence 3029	AF174595	GENBANK	Sequence 30		GENBANK
Sequence 3030	AF176574	GENBANK	Sequence 30		GENBANK
Sequence 3031	AF176702	GENBANK	Sequence 30		GENBANK
Sequence 3032	AF179274	GENBANK	Sequence 30		GENBANK
Sequence 3033	AF188745	GENBANK	Sequence 30	85 D17069	GENBANK
Sequence 3034	AF188746	GENBANK	Sequence 30	86 D17260	GENBANK
Sequence 3035	AF190167	GENBANK	Sequence 30		GENBANK
Sequence 3036	AF191020	GENBANK	Sequence 30		GENBANK
Sequence 3037	AJ000041	GENBANK	Sequence 30	89 D21243	GENBANK
Sequence 3038	AJ001189	GENBANK	Sequence 30	90 <b>D21</b> 260	GENBANK
Sequence 3039	AJ001838	GENBANK	Sequence 30	91 D23672	GENBANK
Sequence 3040	AJ002744	GENBANK	Sequence 30	92 D25274	GENBANK
Sequence 3041	AJ004913	GENBANK	Sequence 30		GENBANK
Sequence 3042	AJ005821	GENBANK	Sequence 30	94 D25542	GENBANK
Sequence 3043	AJ005893	GENBANK	Sequence 30	95 D26362	GENBANK
Sequence 3044	AJ006470	GENBANK	Sequence 30		GENBANK
Sequence 3045	AJ010953	GENBANK	Sequence 30		GENBANK
Sequence 3046	AJ011497	GENBANK	Sequence 30		GENBANK
Sequence 3047	AJ012499	GENBANK	Sequence 30		GENBANK
Sequence 3048	AJ132694	GENBANK	Sequence 31		GENBANK
Sequence 3049	AJ242829	GENBANK	Sequence 31		GENBANK
Sequence 3050	AJ243874	GENBANK	Sequence 31		GENBANK
Sequence 3051	AL049229	GENBANK	Sequence 31		GENBANK
Sequence 3052	AL049784	GENBANK	Sequence 31		GENBANK
Sequence 3053	AL049934	GENBANK	Sequence 31		GENBANK
Sequence 3054	AL050037	GENBANK	Sequence 31		GENBANK
Sequence 3055	AL050101	GENBANK	Sequence 31		GENBANK
Sequence 3056	AL050136	GENBANK	Sequence 31		GENBANK
Sequence 3057	AL050265	GENBANK	Sequence 31		GENBANK
Sequence 3058	AL050274	GENBANK	Sequence 31		GENBANK
Sequence 3059	AL050282	GENBANK	Sequence 31		GENBANK
Sequence 3060	AL050373	GENBANK	Sequence 31		GENBANK
Sequence 3061	AL079279	GENBANK	Sequence 31		GENBANK
Sequence 3062	AL080135	GENBANK	Sequence 31		GENBANK
Sequence 3063	AL080172	GENBANK	Sequence 31		GENBANK
Sequence 3064	AL080192	GENBANK	Sequ <b>ence</b> 31	16 D63997	GENBANK

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Sequence 3	117	D67029	GENBANK	Sequence 3169	L10284	GENBANK
Sequence 3	118	D79991	GENBANK	Sequence 3170	L11566	GENBANK
Sequence 3	119	D79994	GENBANK	Sequence 3171	L12387	GENBANK
Sequence 3	120	D83032	GENBANK	Sequence 3172	L13434	GENBANK
Sequence 3	121	D83703	GENBANK	Sequence 3173	L13852	GENBANK
Sequence 3	122	D84476	GENBANK	Sequence 3174	L15203	GENBANK
Sequence 3	123	D86972	GENBANK	Sequence 3175	L19597	GENBANK
Sequence 3	124	D86984	GENBANK	Sequence 3176	L19713	GENBANK
Sequence 3	125	D87292	GENBANK	Sequence 3177	L20010	GENBANK
Sequence 3	126	D87328	GENBANK	Sequence 3178	L20859	GENBANK
Sequence 3	127	D87437	GENBA <b>NK</b>	Sequence 3179	L20941	GENBANK
Sequence 3	128	D87438	GENBANK	Sequence 3180	L22009	GENBANK
Sequence 3	129	D87452	GENBANK	Sequence 3181	L25081	GENBANK
Sequence 3	130	D87455	GENBANK	Sequence 3182	L25085	GENBANK
Sequence 3	131	D87666	GENBANK	Sequence 3183	L25610	GENBANK
Sequence 3	132	D87667	GENBANK	Sequence 3184	L28010	GENBANK
Sequence 3	133	D87735	GENBANK	Sequence 3185	L29008	GENBANK
Sequence 3	134	D87953	GENBANK	Sequence 3186	L34087	GENBANK
Sequence 3	135	D89289	GENBANK	Sequence 3187	L34839	GENBANK
Sequence 3	136	D89675	GENBANK	Sequence 3188	L38951	GENBANK
Sequence 3	137	D90228	GENBANK	Sequence 3189	L40403	GENBANK
Sequence 3	138	D90427	GENBANK	Sequence 3190	L41143	GENBANK
Sequence 3	139	E01500	GENBANK	Sequence 3191	L42542	GENBANK
Sequence 3	140	E02135	GENBANK	Sequence 3192	L47345	GENBANK
Sequence 3	141	E02628	GENBANK	Sequence 3193	L49399	GENBANK
Sequence 3	142	E06721	GENBANK	Sequence 3194	M10036	GENBANK
Sequence 3	143	J00200	GENBANK	Sequence 3195	M10119	GENBANK
Sequence 3	144	J02871	GENBANK	Sequence 3196	M11233	GENBANK
Sequence 3	145	J03007	GENBANK	Sequence 3197	M11353	GENBANK
Sequence 3	146	J03040	GENBANK	Sequence 3198	M11560	GENBANK
Sequence 3	147	J03464	GENBANK	Sequence 3199	M12938	GENBANK
Sequence 3	148	J03473	GENBANK	Sequence 3200	M13520	GENBANK
Sequence 3	149	J03544	GENBANK	Sequence 3201	M13692	GENBANK
Sequence 3		J03592	GENBANK	Sequence 3202	M15395	GENBANK
Sequence 3		J03779	GENBANK	Sequence 3203	M15796	GENBANK
Sequence 3		J03799	GENBANK	Sequence 3204	M15885	GENBANK
Sequence 3		J03827	GENBANK	Sequence 3205	M15990	GENBANK
Sequence 3		J04177	GENBANK	Sequence 3206	M16247	GENBANK
Sequence 3		J04443	GENBANK	Sequence 3207	M16553	GENBANK
Sequence 3		J04478	GENBANK	Sequence 3208	M17254	GENBANK
Sequence 3		J04615	GENBANK	Sequence 3209	M17885	GENBANK
Sequence 3		J04759	GENBANK	Sequence 3210	M17886	GENBANK
Sequence 3		J05176	GENBANK	Sequence 3211	M20260	GENBANK
Sequence 3		J05192	GENBANK	Sequence 3212	M21895	GENBANK
Sequence 3		J05211	GENBANK	Sequence 3213	M21896	GENBANK
Sequence 3		J05633	GENBANK	Sequence 3214	M22918	GENBANK
Sequence 3		K00409	GENBANK	Sequence 3215	M22920	GENBANK
Sequence 3		K01911	GENBANK	Sequence 3216	M23613	GENBANK
Sequence 3		L01042	GENBANK	Sequence 3217	M24194	GENBANK
Sequence 3		L05092	GENBANK	Sequence 3218	M24594	GENBANK
Sequence 3		L07033	GENBANK	Sequence 3219	M26663	GENBANK
Sequence 3	168	L08044	GENBANK	Sequence 3220	M28016	GENBANK

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Sequence 3221	M28211	GENBANK	Sequence 3273	U09367	GENBANK
Sequence 3222	M29366	GENBANK	Sequen <b>ce 3274</b>	U09410	GENBANK
Sequence 3223	M29536	GENBANK	Sequence 3275	U10248	GENBANK
Sequence 3224	M30448	GENBANK	Sequence 3276	U12465	GENBANK
Sequence 3225	M30818	GENBANK	Sequence 3277	U14603	GENBANK
Sequence 3226	<b>M3</b> 1899	GENBANK	Sequence 3278	U14968	GENBANK
Sequence 3227	M34840	GENBANK	Sequence 3279	U14969	GENBANK
Sequence 3228	M35252	GENBANK	Sequence 3280	U14970	GENBANK
Sequence 3229	M36072	GENBANK	Sequence 3281	U17077	GENBANK
Sequence 3230	M36647	GENBANK	Sequence 3282	U17104	GENBANK
Sequence 3231	M38690	GENBANK	Sequence 3283	U17105	GENBANK
Sequence 3232	M55409	GENBANK	Sequence 3284	U17838	GENBANK
Sequence 3233	<b>M59</b> 849	GENBANK	Sequence 3285	U18197	GENBANK
Sequence 3234	M60457	GENBANK	Sequence 3286	U20770	GENBANK
Sequence 3235	M60828	GENBANK	Sequence 3287	U21090	GENBANK
Sequence 3236	M61831	GENBANK	Sequence 3288	U22897	GENBANK
Sequence 3237	M61866	GENBANK	Sequence 3289	U25766	GENBANK
Sequence 3238	<b>M61</b> 916	GENBANK	Sequence 3290	U29091	GENBANK
Sequence 3239	M63573	GENBANK	Sequence 3291	U30897	GENBANK
Sequence 3240	M64098	GENBANK	Sequence 3292	U33760	GENBANK
Sequence 3241	M64241	GENBANK	Sequence 3293	U34038	GENBANK
Sequence 3242	M64572	GENBANK	Sequence 3294	U34994	GENBANK
Sequence 3243	M68840	GENBANK	Sequence 3295	U37230	GENBANK
Sequence 3244	M68864	GENBANK	Sequence 3296	U37518	GENBANK
Sequence 3245	M69177	GENBANK	Sequence 3297	U37558	GENBANK
Sequence 3246	M69181	GENBANK	Sequence 3298	U37690	GENBANK
Sequence 3247	M73547	GENBANK	Sequence 3299	U39840	GENBANK
Sequence 3248	M74509	GENBANK	Sequence 3300	U39945	GENBANK
Sequence 3249	M77142	GENBANK	Sequence 3301	U40272	GENBANK
Sequence 3250	M77830	GENBANK	Sequence 3302	U40572	GENBANK
Sequence 3251	M80783	GENBANK	Sequence 3303	U43701	GENBANK
Sequence 3252	M81757	GENBANK	Sequence 3304	U44839	GENBANK
Sequence 3253	M83205	GENBANK	Sequence 3305	U46751	GENBANK
Sequence 3254	M84711	GENBANK	· Sequence 3306	U47674	GENBANK
Sequence 3255	M90054	GENBANK	Sequence 3307	U47742	GENBANK
Sequence 3256	M90309	GENBANK	Sequence 3308	<b>U</b> 49 <b>9</b> 57	GENBANK
Sequence 3257	M94345	GENBANK	Sequence 3309	U51134	GENBANK
Sequence 3258	M95627	GENBANK	Sequence 3310	<b>U5</b> 1678	GENBANK
Sequence 3259	S42658	GENBANK	Sequence 3311	U52969	GENBANK
Sequence 3260	S48196	GENBANK	Sequence 3312	U56255	GENBANK
Sequence 3261	S50015	GENBANK	Sequence 3313	U60116	GENBANK
Sequence 3262	S63912	GENBANK	Sequence 3314	U63289	GENBANK
Sequence 3263	S69272	GENBANK	Sequence 3315	U64315	GENBANK
Sequence 3264	S74678	GENBANK	Sequence 3316	U65785	GENBANK
Sequence 3265	S75755	GENBANK	Sequence 3317	<b>U6</b> 6197	GENBANK
Sequence 3266	S76756	GENBANK	Sequence 3318	U66469	GENBANK
Sequence 3267	S78203	GENBANK	Sequence 3319	U67171	GENBANK
Sequence 3268	U00930	GENBANK	Sequence 3320	U68140	GENBANK
Sequence 3269	U <b>01</b> 184	GENBANK	Sequence 3321	U70323	GENBANK
Sequence 3270	U <b>03</b> 688	GENBANK	Sequence 3322	U71363	GENBANK
Sequence 3271	U <b>042</b> 09	GENBANK	Sequence 3323	U73824	GENBANK
Sequence 3272	U08815	GENBANK	Sequence 3324	U75272	GENBANK

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Sequence 3325	U75329	GENBANK	Sequence 3377	X74801	GENBANK
Sequence 3326	U77720	GENBANK	Sequence 3378	X75304	GENBANK
Sequence 3327	U80040	GENBANK	Sequence 3379	X75535	GENBANK
Sequence 3328	U80213	GENBANK	Sequence 3380	X77753	GENBANK
Sequence 3329	U82226	GENBANK	Sequence 3381	X78136	GENBANK
Sequence 3330	U82756	GENBANK	Sequence 3382	X81889	GENBANK
Sequence 3331	U83115	GENBANK	Sequence 3383	X85134	GENBANK
Sequence 3332	U89505	GENBANK	Sequ <b>ence 3384</b>	X87241	GENBANK
Sequence 3333	U90907	GENBANK	Sequ <b>ence 3385</b>	X87949	GENBANK
Sequence 3334	U90942	GENBANK	Sequence 3386	X92689	GENBANK
Sequence 3335	U92458	GENBANK	Sequence 3387	X94754	GENBANK
Sequence 3336	U95825	GENBANK	Sequence 3388	X97335	GENBANK
Sequence 3337	X01037	GENBANK	Sequence 3389	Y00052	GENBANK
Sequence 3338	X01742	GENBANK	Sequence 3390	Y00815	GENBANK
Sequence 3339	X02530	GENBANK	Sequ <b>ence 3391</b>	Y09188	GENBANK
Sequence 3340	X02544	GENBANK	Sequence 3392	Y11164	GENBANK
Sequence 3341	X02596	GENBANK	Sequence 3393	Y12781	GENBANK
Sequence 3342	X04098	GENBANK	Sequence 3394	Y15286	GENBANK
Sequence 3343	X05332	GENBANK	Sequence 3395	Z22968	GENBANK
Sequence 3344	X06272	GENBANK	Sequence 3396	Z36715	GENBANK
Sequence 3345	X06747	GENBANK	Sequence 3397	Z47087	GENBANK
Sequence 3346	X12451	GENBANK	Sequence 3398	Z50749	GENBANK
Sequence 3347	X12654	GENBANK	Sequence 3399	AA001999	dbEST
Sequence 3348	X13238	GENBANK	Sequence 3400	AA004255	dbEST
Sequence 3349	X13425	GENBANK	Sequence 3401	AA004869	dbEST
Sequence 3350	X14420	GENBANK	Sequence 3402	AA007389	dbEST
Sequence 3351	X15653	GENBANK	Sequence 3403	AA007474	dbEST
Sequence 3352	X15729	GENBANK	Sequence 3404	AA009527	dbEST
Sequence 3353	X15822	GENBANK	Sequence 3405	AA010167	dbEST
Sequence 3354	X15949	GENBANK	Sequence 3406	AA010284	dbEST
Sequence 3355	X16940	GENBANK	Sequence 3407	AA011297	dbEST
Sequence 3356	X17206	GENBANK	Sequence 3408	AA011656	dbEST
Sequence 3357	X54941	GENBANK	Sequence 3409	AA011707	dbEST
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Sequence 3359	X56932	GENBANK	Sequence 3411	AA017512	dbEST
Sequence 3360	X57398	GENBANK	Sequence 3412	AA017614	dbEST
Sequence 3361	X57500	GENBANK	Sequence 3413	AA018102	dbEST
Sequence 3362	X58141	GENBANK	Sequence 3414	AA018202	dbEST
Sequence 3363	X60111	GENBANK	Sequence 3415	AA018345	dbEST
Sequence 3364	X63422	GENBANK	Sequence 3416	AA018888	dbEST
Sequence 3365	X63432	GENBANK	Sequence 3417	AA019538	dbEST
Sequence 3366	X64707	GENBANK	Sequence 3418	AA021544	dbEST
Sequence 3367	X65923	GENBANK	Sequence 3419	AA022965	dbEST
Sequence 3368	X68277	GENBANK	Sequence 3420	AA025287	dbEST
Sequence 3369	X68560	GENBANK	Sequence 3421	AA028103	dbEST
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Sequence 3371	X69086	GENBANK	Sequence 3423	AA029143	dbEST
Sequence 3372	X69111	GENBANK	Sequence 3424	AA031367	dbEST
Sequence 3373	X71129	GENBANK	Sequence 3425	AA032194	dbEST
Sequence 3374	X72727	GENBANK	Sequence 3426	AA033869	dbEST
Sequence 3375	X73114	GENBANK	Sequence 3427	AA034094	dbEST
Sequence 3376	X73608	GENBANK	Sequence 3428	AA035549	dbEST

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		"- COT	Sequence 3481	AA082829	dbEST
Sequence 3429	AA035604	dbEST	Sequence 3482	AA083148	dbEST
Sequence 3430	AA035667	dbEST	Sequence 3483	AA083472	dbEST
Sequence 3431	AA035773	dbEST	Sequence 3484	AA083573	dbEST
Sequence 3432	AA037138	dbEST	Sequence 3485	AA083654	dbEST
Sequence 3433	AA037143	dbEST	Sequence 3486	AA083941	dbEST
Sequence 3434	AA041205	dbEST	Sequence 3487	AA085810	dbEST
Sequence 3435	AA041250	dbEST	Sequence 3488	AA086007	dbEST
Sequence 3436	AA043141	dbEST	Sequence 3489	AA086087	dbEST
Sequence 3437	AA044652	dbEST	Sequence 3490	AA088655	dbEST
Sequence 3438	AA044786	dbEST	Sequence 3491	AA088758	dbEST
Sequence 3439	AA045240	dbEST	Sequence 3492	AA088914	dbEST
Sequence 3440	AA045606	dbEST	Sequence 3493	AA089566	dbEST
Sequence 3441	AA045992	dbEST	Sequence 3494	AA090023	dbEST
Sequence 3442	AA047482	dbEST		AA090106	dbEST
Sequence 3443	AA047618	dbEST	Sequence 3495	AA090625	dbEST
Sequence 3444	AA053192	dbEST	Sequence 3496	AA091245	dbEST
Sequence 3445	AA053459	dbEST	Sequence 3497	AA092596	dbEST
Sequence 3446	AA054778	dbEST	Sequence 3498	AA092811	dbEST
Sequence 3447	AA055285	dbEST	Sequence 3499	AA093359	dbEST
Sequence 3448	AA055326	dbEST	Sequence 3500	AA095002	dbEST
Sequence 3449	AA055475	dbEST	Sequence 3501	AA096320	dbEST
Sequence 3450	AA055650	dbEST	Sequence 3502		dbEST
Sequence 3451	AA056025	dbEST	Sequence 3503	AA096453	dbEST
Sequence 3452	AA056369	dbEST	Sequence 3504	AA098824	dbEST
Sequence 3453	AA057721	dbEST	Sequence 3505	AA099424	dbEST
Sequence 3454	AA058458	dbEST	Sequence 3506	AA099718	dbEST
Sequence 3455	AA058488	dbEST	Sequence 3507	AA100133	dbEST
Sequence 3456	AA058899	dbEST	Sequence 3508	AA101276	dbEST
Sequence 3457	AA063476	dbEST	Sequence 3509	AA101476	dbEST
Sequence 3458		dbEST	Sequence 3510	AA102159	dbEST
Sequence 3459		dbEST	Sequence 3511	AA111892	dbEST
Sequence 3460		dbEST	Sequence 3512	AA112162	dbEST
Sequence 3461		dbEST	Sequence 3513		dbEST
Sequence 3462		dbEST	Sequence 3514	AA113952	
Sequence 3463		dbEST	Sequence 3515	AA114003	dbEST
Sequence 3464		dbEST	Sequence 3516	AA114061	dbEST
Sequence 3465		dbEST	Sequence 3517	AA115215	dbEST
Sequence 3468		dbEST	Sequence 3518	AA120849	dbEST
Sequence 3467		dbEST	Sequence 3519		dbEST
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Sequence 3469		dbEST	Sequence 352	1 AA126621	dbEST
Sequence 3470		dbEST	Sequence 3522		dbEST
Sequence 347		dbEST	Sequence 352		_
Sequence 347		dbEST	Sequence 352	4 AA127185	dbEST
Sequence 347		dbEST	Sequence 352	5 AA127436	dbEST
		dbEST	Sequence 352	6 AA127552	dbEST
Sequence 347		dbEST	Sequence 352		dbEST
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Sequence 347		dbEST	Sequence 353	0 AA130615	dbEST
Sequence 347		dbEST	Sequence 353	1 AA131077	dbEST
Sequence 347		dbEST	Sequence 353		dbEST
Sequence 348	0 44002012	40201	•		

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Sequence 3533	AA131233	dbEST	Sequence 3585	AA179187	dbEST
Sequence 3534	AA132802	dbEST	Sequence 3586	AA179362	dbEST
Sequence 3535	AA132857	dbEST	Sequence 3587	AA179567	dbEST
Sequence 3536	AA132862	dbEST	Sequence 3588	AA179691	dbEST
Sequence 3537	AA133071	dbEST	Sequence 3589	AA180398	dbEST
Sequence 3538	AA133248	dbEST	Sequence 3590	AA180914	dbEST
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Sequence 3540	AA133724	dbEST	Sequence 3592	AA181855	dbEST
Sequence 3541	AA133982	dbEST	Sequence 3593	AA182804	dbEST
Sequence 3542	AA134031	dbEST	Sequence 3594	AA182839	dbEST
Sequence 3543	AA134321	dbEST	Sequence 3595	AA186608	dbEST
Sequence 3544	AA134377	dbEST	Sequence 3596	AA187629	dbEST
Sequence 3545	AA135384	dbEST	Sequence 3597	AA188045	dbEST
Sequence 3546	AA135454	dbEST	Sequence 3598	AA188065	dbEST
Sequence 3547	AA135911	dbEST	Sequence 3599	AA188101	dbEST
Sequence 3548	AA136163	dbEST	Sequence 3600	AA188780	dbEST
Sequence 3549	AA143604	dbEST	Sequence 3601	AA188867	dbEST
Sequence 3550	AA146916	dbEST	Sequence 3602	AA190779	dbEST
Sequence 3551	AA147403	dbEST	Sequence 3603	AA190890	dbEST
Sequence 3552	AA147898	dbEST	Sequence 3604	AA191092	dbEST
Sequence 3553	AA147902	dbEST	Sequence 3605	AA191719	dbEST
Sequence 3554	AA148250	dbEST	Sequence 3606	AA192604	dbEST
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Sequence 3557	AA150891	dbEST	Sequence 3609	AA194577	dbEST
Sequence 3558	AA151018	dbEST	Sequence 3610	AA194754	dbEST
Sequence 3559	AA151163	dbEST	Sequence 3611	AA195377	dbEST
Sequence 3560	AA151401	dbEST	Sequence 3612	AA203110	dbEST
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Sequence 3562	AA151979	dbEST	Sequence 3614	AA205928	dbEST
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Sequence 3566	AA157163	dbEST	Sequence 3618	AA206991	dbEST
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Sequence 3575	AA160630	dbEST	Sequence 3627	AA220219	dbES <b>T</b>
Sequence 3576	AA164677	dbEST	Sequence 3628	AA220970	dbEST
Sequence 3577	AA165148	dbEST	Sequence 3629	AA223564	dbEST
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Sequence 3581	AA173331	dbEST	Sequence 3633	AA225294	dbEST
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Sequence 3583	AA176178	dbEST	Sequence 3635	AA226012	dbEST
Sequence 3584	AA179098	dbEST	Sequence 3636	AA226101	dbEST
			,	_	

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Sequence 3637	AA226171	dbEST	Sequence 3689	AA293036	dbEST dbEST
Sequence 3638	AA226430	dbEST	Sequence 3690	AA293047	dbEST
Sequence 3639	AA227059	dbEST	Sequence 3691	AA293050	dbEST
Sequence 3640	AA227549	dbEST	Sequence 3692	AA296186	dbEST
Sequence 3641	AA228012	dbEST	Sequence 3693	AA296326	
Sequence 3642	AA229495	dbEST	Sequence 3694	AA296502	dbEST
Sequence 3643	AA232186	dbEST	Sequence 3695	AA296812	dbEST
Sequence 3644	AA233021	dbEST	Sequence 3696	AA296846	dbEST
Sequence 3645	AA233233	dbEST	Sequence 3697	AA296970	dbEST
Sequence 3646	AA233687	dbEST	Sequence 3698	AA297270	dbEST
Sequence 3647	AA234300	dbEST	Sequence 3699	AA297402	dbEST
Sequence 3648	AA234464	dbEST	Sequence 3700	AA298199	dbEST
Sequence 3649	AA234533	dbEST	Sequence 3701	AA298732	dbEST
Sequence 3650	AA234698	dbEST	Sequence 3702	AA298986	dbEST
Sequence 3651	AA235914	dbEST	Sequence 3703	AA299541	dbEST
Sequence 3652	AA236865	dbEST	Sequence 3704	AA299961	dbEST
Sequence 3653	AA242891	dbEST	Sequence 3705	AA300065	dbEST
Sequence 3654	AA242951	dbEST	Sequence 3706	AA300262	dbEST
Sequence 3655	AA244003	dbEST	Sequence 3707	AA301753	dbEST
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Sequence 3660	AA251262	dbEST	Sequence 3712	AA303334	dbEST
Sequence 3661	AA255576	dbEST	Sequence 3713	AA303594	dbEST
Sequence 3662	AA256248	dbEST	Sequence 3714	AA303657	dbEST
Sequence 3663	AA256403	dbEST	Sequence 3715	AA303696	dbEST
Sequence 3664	AA256724	dbEST	Sequence 3716	AA304025	dbEST
Sequence 3665	AA258750	dbEST	Sequence 3717	AA305121	dbEST
Sequence 3666	AA259025	dbEST	Sequence 3718	AA305231	dbEST
Sequence 3667	AA262494	dbEST	Sequence 3719	AA305315	dbEST
Sequence 3668	AA263002	dbEST	Sequence 3720	AA305333	dbEST
Sequence 3669	AA263040	dbEST	Sequence 3721	AA305566	dbEST
Sequence 3670	AA263145	dbEST	Sequence 3722	AA305895	dbEST
Sequence 3671	AA278642	dbEST	Sequence 3723	AA305951	dbEST
Sequence 3672	AA278729	dbEST	Sequence 3724	AA306028	dbEST
Sequence 3673	AA280080	dbEST	Sequence 3725	AA306129	dbEST
Sequence 3674	AA280089	dbEST	Sequence 3726	AA306402	dbEST
Sequence 3675	AA280876	dbEST	Sequence 3727	AA306668	dbEST
Sequence 3676	AA282734	dbEST	Sequence 3728	AA306983	dbEST
Sequence 3677	AA283746	dbEST	Sequence 3729	AA307101	dbEST
Sequence 3678	AA284642	dbEST	Sequence 3730	AA307579	dbEST
Sequence 3679	AA284806	dbEST	Sequence 3731	AA307590	dbEST
Sequence 3680		dbEST	Sequence 3732	AA307728	dbEST
Sequence 3681	AA286841	dbEST	Sequence 3733	AA307746	dbEST
Sequence 3682		dbEST	Sequence 3734	AA307789	dbEST
Sequence 3683		dbEST	Sequence 3735	AA307941	dbEST
Sequence 3684		dbEST	Sequence 3736		dbEST
Sequence 3685		dbEST	Sequence 3737	AA308269	dbEST
Sequence 3686		dbEST	Sequence 3738		dbEST
Sequence 3687		dbEST	Sequence 3739		dbEST
•		dbEST	Sequence 3740		dbEST
Sequence 3688	MY727130	aneo:	<b>⇒=</b> ¶===		

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Sequence 3741	AA309445	dbEST	Sequence 3793	AA318 <b>544</b>	dbEST
Sequence 3742	AA309812	dbEST	Sequence 3794	AA319139	dbEST
Sequence 3743	AA3101 <b>63</b>	dbEST	Sequence 3795	AA319538	dbEST
Sequence 3744	AA310181	dbEST	Sequence 3796	AA320701	dbEST
Sequence 3745	AA3102 <b>92</b>	dbEST	Sequence 3797	AA322246	dbEST
Sequence 3746	AA3104 <b>60</b>	dbEST	Sequence 3798	AA324251	dbEST
Sequence 3747	AA3105 <b>61</b>	dbEST	Sequence 3799	AA324411	dbEST
Sequence 3748	AA310591	dbEST	Sequence 3800	AA324705	dbEST
Sequence 3749	AA310928	dbEST	Sequence 3801	AA324831	dbEST
Sequence 3750	AA311044	dbEST	Sequence 3802	AA325221	dbEST
Sequence 3751	AA311227	dbEST	Sequence 3803	AA325658	dbEST
Sequence 3752	AA312177	dbEST	Sequence 3804	AA326793	dbEST
Sequence 3753	AA312242	dbEST	Sequence 3805	AA327228	dbEST
Sequence 3754	AA312444	dbEST	Sequence 3806	AA327546	dbEST
Sequence 3755	AA312448	dbEST	Sequence 3807	AA327611	dbEST
Sequence 3756	AA312503	dbEST	Sequence 3808	AA330234	dbEST
Sequence 3757	AA313200	dbEST	Sequence 3809	AA330784	dbEST
Sequence 3758	AA313208	dbEST	Sequence 3810	AA331859	dbEST
Sequence 3759	AA313418	dbEST	Sequence 3811	AA332556	dbEST
Sequence 3760	AA313549	dbEST	Sequence 3812	AA332667	dbEST
Sequence 3761	AA313828	dbEST	Sequence 3813	AA333307	doEST
Sequence 3762	AA313837	dbEST	Sequence 3814	AA333526	doEST
Sequence 3763	AA313933	dbEST	Sequence 3815	AA334452	dbEST
Sequence 3764	AA313996	dbEST	Sequence 3816	AA336081	dbEST
Sequence 3765	AA314061	dbEST	Sequence 3817	AA336683	dbEST
Sequence 3766	AA314181	dbEST	Sequence 3818	AA336740	dbEST
Sequence 3767	AA314188	dbEST	Sequence 3819	AA338257	dbEST
Sequence 3768	AA314202	dbEST	Sequence 3820	AA339065	dbEST
Sequence 3769	AA314241	dbEST	Sequence 3821	AA3391 <b>51</b>	dbEST
Sequence 3770	AA314355	dbEST	Sequence 3822	AA340654	dbEST
Sequence 3771	AA314465	dbEST	Sequence 3823	AA340719	dbEST
Sequence 3772	AA314748	dbEST	Sequence 3824	AA344393	dbEST
Sequence 3773	AA3147 <b>57</b>	dbEST	Sequence 3825	AA345558	dbEST
Sequence 3774	AA314847	dbEST	Sequence 3826	AA346918	dbEST
Sequence 3775	AA314862	dbEST	Sequence 3827	AA347293	dbEST
Sequence 3776	AA3148 <b>82</b>	dbEST	Sequence 3828	AA347390	dbEST
Sequence 3777	AA314961	dbEST	Sequence <b>382</b> 9	AA3478 <b>8</b> 9	dbEST
Sequence 3778	AA315069	dbEST	Sequence 3830	AA349118	dbEST
Sequence 3779	AA315172	dbEST	Sequence 3831	AA350206	dbEST
Sequence 3780	AA31521 <b>5</b>	dbEST	Sequence 3832	AA350228	dbEST
Sequence 3781	AA315321	dbEST	Sequence 3833	AA350661	dbEST
Sequence 3782	AA315406	dbEST	Sequence 3834	AA352755	dbEST
Sequence 3783	AA315444	dbEST	Sequence 3835	AA354391	dbEST
Sequence 3784	AA315497	dbEST	Sequence 3836	AA355003	dbEST
Sequence 3785	AA315762	dbEST	Sequence 3837	AA355196	dbEST
Sequence 3786	AA3159 <b>50</b>	dbEST	Sequence 3838	AA355363	dbEST
Sequence 3787	AA3159 <b>93</b>	dbEST	Sequence 3839	AA355451	dbEST
Sequence 3788	AA316199	dbEST	Sequence 3840	AA359003	dbEST
Sequence 3789	AA316322	dbEST	Sequence 3841	AA3592 <b>84</b>	dbEST
Sequence 3790	AA317096	dbEST	Sequence 3842	AA359514	dbEST
Sequence 3791	AA317937	dbEST	Sequence 3843	AA361042	dbEST
Sequence 3792	AA3179 <b>56</b>	dbEST	Sequence 3844	AA362778	dbEST

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Sequence 3845	AA362825	dbEST	Sequence 3897	AA425613	dbEST
Sequence 3846	AA363142	dbEST	Sequence 3898	AA426235	dbEST
Sequence 3847	AA364498	dbEST	Sequence 3899	AA426391	dbEST
Sequence 3848	AA365150	dbEST	Sequence 3900	AA427471	dbEST
Sequence 3849	AA366518	dbEST	Sequence 3901	AA427596	dbEST
Sequence 3850	AA367581	dbEST	Sequence 3902	AA428096	dbEST
Sequence 3851	AA368810	dbEST	Sequence 3903	AA429173	dbEST
Sequence 3852	AA371017	dbEST	Sequence 3904	AA430304	dbEST
Sequence 3853	AA371265	dbEST	Sequence 3905	AA430478	dbEST
Sequence 3854	AA373756	dbEST	Sequence 3906	AA430597	dbEST
Sequence 3855	AA374222	dbEST	Sequence 3907	AA430628	dbEST
Sequence 3856	AA374766	dbEST	Sequence 3908	AA43402B	dbEST
Sequence 3857	AA375483	dbEST	Sequence 3909	AA436008	dbEST
Sequence 3858	AA377218	dbEST	Sequence 3910	AA436762	dbEST
	AA378059	dbEST	Sequence 3911	AA436769	dbEST
Sequence 3859	AA378226	dbEST	Sequence 3912	AA437277	dbEST
Sequence 3860	AA379015	dbEST	Sequence 3913	AA442287	dbEST
Sequence 3861	AA385388	dbEST	Sequence 3914	AA442517	dbEST
Sequence 3862	AA386267	dbEST	Sequence 3915	AA442687	dbEST
Sequence 3863	AA393259	dbEST	Sequence 3916	AA443659	dbEST
Sequence 3864	AA393906	dbEST	Sequence 3917	AA446071	dbEST
Sequence 3865	AA394176	dbEST	Sequence 3918	AA446716	dbEST
Sequence 3866	AA394299	dbEST	Sequence 3919	AA446958	dbEST
Sequence 3867	AA397802	dbEST	Sequence 3920	AA447885	dbEST
Sequence 3868	AA399239	dbEST	Sequence 3921	AA448118	dbEST
Sequence 3869	AA399393	dbEST	Sequence 3922	AA448950	dbEST
Sequence 3870	AA399637	dbEST	Sequence 3923	AA450168	dbEST
Sequence 3871	AA401603	dbEST	Sequence 3924	AA451733	dbEST
Sequence 3872	AA401957	dbEST	Sequence 3925	AA452012	dbEST
Sequence 3873	AA402792	dbEST	Sequence 3926	AA453445	dbEST
Sequence 3874	AA403235	dbEST	Sequence 3927	AA453562	dbEST
Sequence 3875	AA403233 AA403322	dbEST	Sequence 3928	AA454027	dbEST
Sequence 3876	AA404444	dbEST	Sequence 3929	AA454065	dbEST
Sequence 3877	AA404642	dbEST	Sequence 3930	AA454912	dbEST
Sequence 3878 Sequence 3879	AA404646	dbEST	Sequence 3931	AA455299	dbEST
Sequence 3880	AA405640	dbEST	Sequence 3932	AA456372	dbEST
Sequence 3881	AA406402	dbEST	Sequence 3933	AA457021	dbEST
Sequence 3882	AA410700	dbEST	Sequence 3934	AA458804	dbEST
Sequence 3883	AA411201	dbEST	Sequence 3935	AA459528	dbEST
Sequence 3884	AA416781	dbEST	Sequence 3936	AA460511	dbEST
Sequence 3885	AA418473	dbEST	Sequence 3937	AA460755	dbEST
Sequence 3886	AA420523	dbEST	Sequence 3938	AA4609 <b>69</b>	dbEST
Sequence 3887	AA420647	dbEST	Sequence 3939	AA464763	dbEST
Sequence 3888		dbEST	Sequence 3940	AA465494	dbEST
Sequence 3889		dbEST	Sequence 3941	AA4656 <b>97</b>	dbEST
Sequence 3890		dbEST	Sequence 3942	AA467864	dbEST
Sequence 3891	AA421213	dbEST	Sequence 3943	AA4679 <b>74</b>	dbEST
•		dbEST	Sequence 3944	AA468839	dbEST
Sequence 3892		dbEST	Sequence 3945	AA469129	dbEST
Sequence 3893		dbEST	Sequence 3946		dbEST
Sequence 3894		dbEST	Sequence 3947	AA4693 <b>0</b> 4	dbEST
Sequence 3895		dbEST	Sequence 3948		dbEST
Sequence 3896	AV-1424000	UDEO I	50400	-	

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Sequence		AA471070	dbEST	Se	quence 4001	AA522473	dbEST
Sequen <b>ce</b>	3950	AA476207	dbEST	Se	quence 4002	AA523006	dbEST
Sequence	3951	AA476543	dbEST	Se	quence 4003	AA523902	dbEST
Sequence	3952	AA476648	dbEST	Se	quence 4004	AA524258	dbEST
Sequence	3953	AA476679	dbEST	Se	quence 4005	AA525497	dbEST
Sequence	3954	AA477579	dbEST	Se	quence 4006	AA525801	dbEST
Sequence	3955	AA477848	dbEST	Se	quence 4007	AA526187	dbEST
Sequence	3956	AA478298	dbEST	Se	quence 4008	AA526472	dbEST
Sequence	3957	AA478522	dbEST		quence 4009	AA526894	dbEST
Sequence	3958	AA478647	dbEST	Se	quence 4010	AA528202	dbEST
Sequence	3959	AA479287	dbEST	Se	quence 4011	AA531561	dbEST
Sequence	3960	AA479809	dbEST	Se	quence 4012	AA532461	dbEST
Sequence	3961	AA480051	dbEST	Se	quence 4013	AA532852	dbEST
Sequence	3962	AA482522	dbEST		quence 4014	AA533001	dbEST
Sequence	3963	AA482884	dbEST		quence 4015	AA533940	dbEST
Sequence	3964	AA484104	dbEST		quence 4016	AA534419	dbEST
Sequence	3965	AA484284	dbEST		quence 4017	AA534543	dbEST
Sequence		AA484568	dbEST		quence 4018	AA535610	dbEST
Sequence	3967	AA484752	dbEST		quence 4019	AA535695	dbEST
Sequence	3968	AA485853	dbEST		quence 4020	AA535837	dbEST
Sequence	3969	AA486838	dbEST		quence 4021	AA536175	dbEST
Sequence	3970	AA488014	dbEST		quence 4022	AA548238	dbEST
Sequence	3971	AA488592	dbEST	Sec	quence 4023	AA548600	dbEST
Sequence	3972	AA488954	dbEST	Sec	quence 4024	AA551065	dbEST
Sequence	3973	AA489124	dbEST	Sec	quence 4025	AA551252	dbEST
Sequence	3974	AA489160	dbEST		quence 4026	AA551773	dbEST
Sequence	3975	AA489323	dbEST	Sec	quence 4027	AA552154	dbEST
Sequence	3976	AA491983	dbEST	Sec	quence 4028	AA552253	dbEST
Sequence	3977	AA492042	dbEST	Sec	quence 4029	AA552570	dbEST
Sequence	3978	AA492256	dbEST	Sed	quence 4030	AA555160	dbEST
Sequence	3979	AA494295	dbEST	Sed	quence 4031	AA557888	dbEST
Sequence	3980	AA494489	dbEST	Sed	quence 4032	AA558976	dbEST
Sequence	3981	AA503040	dbEST	Sec	quence 4033	AA564005	dbEST
Sequence	3982	AA503330	dbEST	Sec	quence 4034	AA564017	dbEST
Sequence	3983	AA503943	dbEST	Sec	quence 4035	AA565044	dbEST
Sequence	3984	AA504703	dbEST	Sec	quence 4036	AA565448	dbEST
Sequence	3985	AA504969	dbEST	Sed	quence 4037	AA569247	dbEST
Sequence	3986	AA505327	db <b>EST</b>	Sed	quence 4038	AA569354	dbEST ·
Sequence		AA505568	dbEST	Sec	quence 4039	AA569439	dbEST
Sequence		AA506299	dbEST		quence 4040	AA570171	dbEST
Sequence		AA506304	dbEST	Sed	quence 4041	AA570182	dbEST
Sequence		AA506459	dbEST		quence 4042	AA572808	dbEST
Sequence		AA507065	dbEST	Sec	quence 4043	AA572938	cbEST
Sequence		AA507201	dbEST		quence 4044	AA573557	cbEST
Sequence		AA507217	dbEST	Sec	quence 4045	AA573559	abEST
Sequence		AA507472	dbEST		quence 4046	AA573575	dbEST
Sequence		AA507595	dbEST		quence 4047	AA573737	cbEST
Sequence		AA508861	dbEST		quence 4048	AA573762	dbEST
Sequence		AA513597	dbEST	Sec	quence 4049	AA573787	dbEST
Sequence		AA514775	dbEST		quence 4050	AA573811	dbEST
Sequence		AA515132	dbEST		quence 4051	AA573817	dbEST
Sequence	4000	AA516441	dbEST	Sec	quence 4052	AA573910	db <b>EST</b>

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	_		0	A A C C C E 4 4	dbEST
Sequence 4053	AA576252	dbES <b>T</b>	Sequence 4105	AA669544	dbEST
Sequence 4054	AA576404	dbEST	Sequence 4106	AA675892	dbEST
Sequence 4055	AA578265	dbEST	Sequence 4107	AA678873	
Sequence 4056	AA578773	dbEST	Sequence 4108	AA683546	dbEST
Sequence 4057	AA5789 <b>04</b>	dbEST	Sequence 4109	AA687733	dbEST
Sequence 4058	AA578976	dbEST	Sequence 4110	AA693922	dbEST
Sequence 4059	AA579 <b>596</b>	dbEST	Sequence 4111	AA694186	dbEST
Sequence 4060	AA581523	dbEST	Sequence 4112	AA694477	dbEST
Sequence 4061	AA582489	dbEST	Sequence 4113	AA700876	dbEST
Sequence 4062	AA582728	dbEST	Sequence 4114	AA703004	dbEST
Sequence 4063	AA582805	dbEST	Sequence 4115	AA703925	dbEST
Sequence 4064	AA583801	dbEST	Sequence 4116	AA704690	dbEST
Sequence 4065	AA586744	dbEST	Sequence 4117	AA705054	dbEST
Sequence 4066	AA588083	dbEST	Sequence 4118	AA70679 <b>7</b>	dbEST
Sequence 4067	AA588358	dbEST	Sequence 4119	AA709070	dbEST
Sequence 4068	AA588801	dbEST	Sequence 4120	AA713760	dbEST
Sequence 4069	AA592947	dbEST	Sequence 4121	AA714902	dbEST
Sequence 4070	AA593385	dbEST	Sequence 4122	AA715036	dbEST
Sequence 4071	AA593879	dbEST	Sequence 4123	AA719530	dbEST
Sequence 4072	AA595193	dbEST	Sequence 4124	AA721298	dbEST
Sequence 4073	AA595471	dbEST	Sequence 4125	AA722090	dbEST
Sequence 4074	AA595582	dbEST	Sequence 4126	AA744876	dbEST
Sequence 4075	AA598598	dbEST	Sequence 4127	AA757282	dbEST
Sequence 4076	AA599229	dbEST	Sequence 4128	AA758023	dbEST
Sequence 4077	AA599650	dbEST	Sequence 4129	AA761562	dbEST
Sequence 4078	AA602472	dbEST	Sequence 4130	AA761831	dbEST
Sequence 4079	AA602773	dbEST	Sequence 4131	AA769102	dbEST
Sequence 4080	AA602979	dbEST	Sequence 4132	AA769142	dbEST
Sequence 4081	AA603572	dbEST	Sequence 4133	AA770690	dbEST
Sequence 4082	AA610734	dbEST	Sequence 4134	AA772736	dbEST
Sequence 4083	AA614274	dbEST	Sequence 4135	AA775824	dbEST
Sequence 4084	AA620334	dbEST	Sequence 4136	AA776709	dbEST
Sequence 4085	AA630382	dbEST	Sequence 4137	AA776740	dbEST
Sequence 4086	AA631152	dbEST	Sequence 4138	AA777265	dbEST
Sequence 4087	AA632954	dbEST	Sequence 4139	AA777489	dbEST
Sequence 4088	AA633399	dbEST	Sequence 4140	AA778571	dbEST
Sequence 4089	AA633909	dbEST	Sequence 4141	AA804879	dbEST
Sequence 4090	AA640458	dbEST	Sequence 4142	AA808146	dbEST
Sequence 4091	AA640716	dbEST	Sequence 4143	AA808466	dbEST
Sequence 4092	AA641074	dbEST	Sequence 4144	AA813108	dbEST
Sequence 4093	AA644678	dbEST	Sequence 4145	AA813287	dbEST
Sequence 4094	AA648209	dbEST	Sequence 4146	AA814412	dbEST
Sequence 4095	AA649298	dbEST	Sequence 4147	AA814634	dbEST
Sequence 4096	AA650241	dbEST	Sequence 4148	AA815173	dbEST
Sequence 4097	AA654709	dbEST	Sequence 4149	AA824395	dbEST
Sequence 4098	AA658207	dbEST	Sequence 4150	AA826491	dbEST
Sequence 4099	AA659179	dbEST	Sequence 4151	AA832480	dbEST
Sequence 4100	AA659725	dbEST	Sequence 4152	AA835490	dbEST
Sequence 4101	AA661464	dbEST	Sequence 4153	AA836233	dbEST
Sequence 4101	AA661927	dbEST	Sequence 4154	AA838431	dbEST
Sequence 4102	AA664482	dbEST	Sèquence 4155	AA843216	dbEST
Sequence 4103	AA669484	dbEST	Sequence 4156	AA843683	dbEST
Dequence 4 104	, 0 1000-10-1	35E01	234222 7.400		

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Sequence 4157	AA844081	dbEST	Sequence 4209	AI039301	dbEST
Sequence 4158	AA844230	dbEST	Sequence 4210	Al039813	dbEST
Sequence 4159	AA854305	dbEST	Sequence 4211	AI040339	dbEST
Sequence 4160	AA856914	dbEST	Sequence 4212	AI042002	dbEST
Sequence 4161	AA864975	dbEST	Sequence 4213	AI049971	dbEST
Sequence 4162	AA868337	dbEST	Sequence 4214	AI050826	dbEST
Sequence 4163	AA873459	dbEST	Sequence 4215	AI051957	dbEST
Sequence 4164	AA877796	dbEST	Sequence 4216	AI052061	dbEST
Sequence 4165	AA877864	dbEST	Sequence 4217	AI061610	dbEST
Sequence 4166	AA883132	dbEST	Sequence 4218	Al064931	dbEST
Sequence 4167	AA883580	dbEST	Sequence 4219	AI075041	dbEST
Sequence 4168	AA886497	dbEST	Sequence 4220	AI077488	dbEST
Sequence 4169	AA896959	dbEST	Sequence 4221	AI077719	dbEST
Sequence 4170	AA902256	dbEST	Sequence 4222	Al078217	dbEST
Sequence 4171	AA907068	dbEST	Sequence 4223	AI081133	dbEST
Sequence 4172	AA926992	dbEST	Sequence 4224	Al081379	dbEST
Sequence 4173	AA932059	dbEST	Sequence 4225	Al081945	dbEST
Sequence 4174	AA933573	dbEST	Sequence 4226	AI082235	dbEST
Sequence 4175	AA936089	dbEST	Sequence 4227	AI085263	dbEST
Sequence 4176	AA938156	dbEST	Sequence 4228	Al085381	JbEST
Sequence 4177	AA947441	dbEST	Sequence 4229	A1087080	dbEST
Sequence 4178	AA948037	dbEST	Sequence 4230	AI088914	dbEST
Sequence 4179	AA948223	dbEST	Sequence 4231	AI089875	dbEST
Sequence 4180	AA954079	dbEST	Sequence 4232	AI090012	dbEST
Sequence 4181	AA962587	dbEST	Sequence 4233	AI090827	dbEST
Sequence 4182	AA969613	dbEST	Sequence 4234	Al091136	dbEST
Sequence 4183	AA970266	dbEST	Sequence 4235	AI091516	dbEST
Sequence 4184	AA970314	dbEST	Sequence 4236	AI091814	dbEST
Sequence 4185	AA970570	dbEST	Sequence 4237	AI094634	dbEST
Sequence 4186	AA977406	dbEST	Sequence 4238	AI097214	dbEST
Sequence 4187	AA978126	dbEST	Sequence 4239	Al114433	dbEST
Sequence 4188	AA983832	dbEST	Sequence 4240	AI114651	dbEST
Sequence 4189	AA984173	dbEST	Sequence 4241	Al125166	dbEST
Sequence 4190	AA984744	dbEST	Sequence 4242	Al126653	dbEST
Sequence 4191	AA987442	dbEST	Sequence 4243	Al127526	dbEST
Sequence 4192	AA988520	dbEST	Sequence 4244	Al127556	dbEST
Sequence 4193	AB005621	dbEST	Sequence 4245	Al128883	dbEST
Sequence 4194	AF017688	dbEST	Sequence 4246	Al128949	dbEST
Sequence 4195	AF062725	dbEST	Sequence 4247	Al131444	dbEST
Sequence 4196	AF062729	dbEST	Sequence 4248	Al133208	dbEST
Sequence 4197	Al004596	dbEST	Sequence 4249	Al139036	dbEST
Sequence 4198	Al015574	dbEST	Sequence 4250	Al139394	dbEST
Sequence 4199	AI016169	dbEST	Sequence 4251	Al140096	dbEST
Sequence 4200	AI017022	dbEST	Sequence 4252	AI140210	dbEST
Sequence 4201	Al017959	dbEST	Sequence 4253	Al141189	dbEST
Sequence 4202	Al018617	dbEST	Sequence 4254	Al141583	dbEST
Sequence 4203	Al024907	dbEST	Sequence 4255	AI141732	dbEST
Sequence 4204	Al025376	dbEST	Sequence 4256	AI148251	dbEST
Sequence 4205	Al027516	dbEST	Sequence 4257	AI150619	dbEST
Sequence 4206	Al034024	dbEST	Sequence 4258	Al160426	dbEST
Sequence 4207	AI038048	dbEST	Sequence 4259	AI161253	dbEST
Sequence 4208	AI039227	dbEST	Sequence 4260	AI167801	dbEST
			·		

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			~		
Sequence 4261	Al184320	dbEST	Sequence 4313	AI267677	dbEST
Sequence 4262	AI189339	dbEST	Sequence 4314	AI267808	dbEST
Sequence 4263	Al189345	dbEST	Sequence 4315	Al267941	dbEST
Sequence 4264	AI189386	dbEST	Sequence 4316	Al269883	dbEST
Sequence 4265	Al189894	dbEST	Sequence 4317	Al271740	dbEST
Sequence 4266	Al190648	dbEST	Sequence 4318	Al274786	dbEST
Sequence 4267	Al192780	dbEST	Sequence 4319	Al275114	dbEST
Sequence 4268	Al192784	db <b>EST</b>	Sequence 4320	Al275175	dbEST
Sequence 4269	AI193675	dbEST	Sequence 4321	AI277434	dbEST
Sequence 4270	Al198311	dbEST	Sequence 4322	AI277739	dbEST
Sequence 4271	Al198771	dbEST	Sequence 4323	Al279141	dbEST
Sequence 4272	Al198986	dbEST	Sequence 4324	AI281882	dbEST
Sequence 4273	Al199641	dbEST	Sequence 4325	Al284011	dbEST
Sequence 4274	AI199681	dbEST	Sequence 4326	AI287972	dbEST
Sequence 4275	Al199773	dbEST	Sequence 4327	AI288388	dbEST
Sequence 4276	Al200324	dbEST	Sequence 4328	AI288730	dbEST
Sequence 4277	AI200444	dbEST	Sequence 4329	AI288929	dbEST
Sequence 4278	AI201261	dbEST	Sequence 4330	Al289120	dbEST
Sequence 4279	AI203141	dbEST	Sequence 4331	Al291583	dbEST
Sequence 4280	AI203350	dbEST	Sequence 4332	Al292299	dbEST
Sequence 4281	Al207650	dbEST	Sequence 4333	Al306464	dbEST
Sequence 4282	AI214043	dbEST	Sequence 4334	Al309401	dbEST
Sequence 4283	Al216966	dbEST	Sequence 4335	Al312049	dbEST
Sequence 4284	Al216986	dbEST	Sequence 4336	Al332904	dbEST
Sequence 4285	Al216988	dbEST	Sequence 4337	Al333548	dbEST
Sequence 4286	AI217003	dbEST	Sequence 4338	Al335269	dbEST
Sequence 4287	Al217009	dbEST	Sequence 4339	Al340246	dbEST
Sequence 4288	AI217021	dbEST	Sequence 4340	Al341076	dbEST
Sequence 4289	Al219174	dbEST	Sequence 4341	Al341334	dbEST
Sequence 4290	Al244335	dbEST	Sequence 4342	Al341352	dbEST
Sequence 4291	Al246167	dbEST	Sequence 4343	Al343091	dbEST
Sequence 4292	Al248514	dbEST	Sequence 4344	Al343604	dbEST
Sequence 4293	Al253288	dbEST	Sequence 4345	Al346657	dbEST
Sequence 4294	Al253300	dbEST	Sequence 4346	Al346809	dbEST
Sequence 4295	Al253304	dbEST	Sequence 4347	Al347155	dbEST
Sequence 4296	Al253319	dbEST	Sequence 4348	Al349452	dbEST
Sequence 4297	A1253330	dbEST	Sequence 4349	AI350388	dbEST
Sequence 4298	Al253335	dbEST	Sequence 4350	Al352032	dbEST
Sequence 4299	Al253367	dbEST	Sequence 4351	Al356718	dbEST
Sequence 4300	Al253436	dbEST	Sequence 4352	A1360870	dbEST
Sequence 4301	Al261763	dbEST	Sequence 4353	Al361965	dbEST
Sequence 4302	Al262139	dbEST	Sequence 4354	Al362490	dbEST
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Sequence 4304	Al267162	dbEST	Sequence 4356	AI366365	dbEST
Sequence 4305	Al267185	dbEST	Sequence 4357	A1366376	dbEST
Sequence 4306	Al267282	dbEST	Sequence 4358	Al372599	dbEST
Sequence 4307	Al267307	dbEST	Sequence 4359	AI373323	dbEST
Sequence 4308	Al267397	dbEST	Sequence 4360	Al375465	dbEST
Sequence 4309	Al267502	dbEST	Sequence 4361	AI381996	dbEST
Sequence 4310	AI267574	dbEST	Sequence 4362	Al394173	dbEST
Sequence 4311	A1267596	dbE <b>ST</b>	Sequence 4363	A1394498	dbEST
Sequence 4312	Al267652	dbEST	Sequence 4364	AI417833	dbEST

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Sequence 4365	Al421412	dbEST	Sequence 4417	Al684039	dbEST
Sequence 4366	AI432603	dbEST	Sequence 4418	A1684170	dbEST
Sequence 4367	AI434025	dbEST	Sequence 4419	A1686957	dbEST
Sequence 4368	AI434639	dbEST	Sequence 4420	AI690236	dbEST
Sequence 4369	Al436714	dbEST	Sequence 4421	Al692940	dbEST
Sequence 4370	Al439173	dbEST	Sequence 4422	AI692995	dbEST
Sequence 4371	AJ453358	dbEST	Sequence 4423	Al694572	dbEST
Sequence 4372	AJ473779	dbEST	Sequence 4424	A1697815	dbEST
Sequence 4373	Al476254	dbEST	Sequence 4425	Al701119	dbEST
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Sequence 4375	Al499439	dbEST	Sequence 4427	AI732541	dbEST
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Sequence 4377	Al521270	dbEST	Sequence 4429	A1744451	dbEST
Sequence 4378	AI525426	dbEST	Sequence 4430	A1745280	dbEST
Sequence 4379	AI525773	dbEST	Sequence 4431	AI751583	dbEST
Sequence 4380	AI525843	dbEST	Sequence 4432	AI751914	dbEST
Sequence 4381	AI537498	dbEST	Sequence 4433	AI752822	dbEST
Sequence 4382	AI538681	dbEST	Sequence 4434	AI753269	dbEST
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Sequence 4384	A1547309	dbEST	Sequence 4436	A1755085	dbEST
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Sequence 4386	AI557495	dbEST	Sequence 4438	AI761728	dbEST
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Sequence 4394	Al587118	dbEST	Sequence 4446	Al815449	dbEST
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Sequence 4397	A1608968	dbEST	Sequence 4449	AI815966	dbEST
Sequence 4398	Al628609	dbEST	Sequence 4450	AI815976	dbEST
Sequence 4399	Al630282	dbEST	Sequence 4451	AI816033	dbEST
Sequence 4400	Al636577	dbEST	Sequence 4452	AI820995	dbEST
Sequence 4401	Al636911	dbEST	Sequence 4453	A1829303	dbEST
Sequence 4402	Al638327	dbEST	Sequence 4454	A1858066	dbEST
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Sequence 4404	Al651069	dbEST	Sequence 4456	AI866262	dbEST
Sequence 4405	Al651974	dbEST	Sequence 4457	AJ871477	dbEST
Sequence 4406	AI651976	dbEST	Sequence 4458	AI872085	dbEST
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Sequence 4408	Al652555	dbEST	Sequence 4460	AJ879010	dbEST
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Sequence 4412	AJ672362	dbEST	Sequence 4464	AI885833	dbEST
Sequence 4413	AI674583	dbEST	Sequence 4465	AI886418	dbEST
Sequence 4414	Al675889	dbEST	Sequence 4466	AI887517	dbEST
Sequence 4415	Al679629	dbEST	Sequence 4467	Al887664	dbEST
Sequence 4416	Al683183	dbEST	Sequence 4468	AI911703	dbEST

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Sequence 4469	Al914133	dbEST	Sequence 4521	AW026680	dbEST
Sequence 4470	Al917649	dbEST	Sequence 4522	AW027007	dbEST
Sequence 4471	A1924598	dbEST	Sequence 4523	AW027219	dbEST
Sequence 4472	Al929113	dbEST	Sequence 4524	AW029146	dbEST
Sequence 4473	Al929453	db <b>EST</b>	Sequence 4525	AW058559	dbEST
Sequence 4474	Al934153	dbEST	Sequence 4526	AW062376	dbEST
Sequence 4475	Al934595	dbEST	Sequence 4527	AW071906	dbEST
Sequence 4476	Al935291	dbEST	Sequence 4528	AW075433	dbEST
Sequence 4477	Al935816	dbEST	Sequence 4529	AW084950	dbEST
Sequence 4478	AI937060	dbEST	Sequence 4530	AW129510	dbEST
Sequence 4479	AI951021	dbEST	Sequence 4531	AW132009	dbEST
Sequence 4480	AI951536	dbEST	Sequence 4532	AW134746	dbEST
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Sequence 4484	AI973218	dbEST	Sequence 4536	AW160669	dbEST
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Sequence 4488	AL035985	db <b>EST</b>	Sequence 4540	AW163441	dbEST
Sequence 4489	AL036026	dbEST	Sequence 4541	AW167481	dbEST
Sequence 4490	AL036415	dbEST	Sequence 4542	AW169941	dbEST
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Sequence 4493	AL036764	dbEST	Sequence 4545	AW177936	dbEST
Sequence 4494	AL037724	dbEST	Sequence 4546	AW177948	dbEST
Sequence 4495	AL038058	dbEST	Sequence 4547	AW177957	dbEST
Sequence 4496	AL038660	dbEST	Sequence 4548	AW179006	dbEST
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Sequence 4498	AL039517	dbEST	Sequence 4550	AW183601	dbEST
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Sequence 4500	AL040505	dbE <b>ST</b>	Sequence 4552	C02912	dbEST
Sequence 4501	AL040993	dbEST	Sequence 4553	C16620	dbEST
Sequence 4502	AL041470	dbEST	Sequence 4554	C18148	dbEST
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Sequence 4504	AL043245	dbEST	Sequence 4556	C18 <b>57</b> 0	dbEST
Sequence 4505	AL044016	dbEST	Sequence 4557	D58 <b>694</b>	dbEST
Sequence 4506	AL044350	db <b>EST</b>	Sequence 4558	F01327	dbEST
Sequence 4507	AL045492	db <b>EST</b>	Sequence 4559	F05894	dbEST
Sequence 4508	AL046107	dbEST	Sequence 4560	F07432	dbEST
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Sequence 4516	AL120807	dbEST	Sequence 4568	H01419	dbEST
Sequence 4517	AW006392	dbE <b>ST</b>	Sequence 4569	H01979	dbEST
Sequence 4518	AW01 1990	dbE <b>ST</b>	Sequence 4570	H02725	dbEST
Sequence 4519	AWC* 11	dbE <b>ST</b>	Sequence 4571	H06040	dbEST
Sequence 4520	AW026543	dbEST	Sequence 4572	H06313	dbEST
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Sequence 4573	H08929	dbEST	Sequence 4625	R70116	dbEST
Sequence 4574	H10932	dbEST	Sequence 4626	R77854	dbEST
Sequence 4575	H12090	dbEST	Sequence 4627	R85291	dbEST
Sequence 4576	H12254	dbEST	Sequence 4628	T30604	dbEST
Sequence 4577	H17195	dbEST	Sequence 4629	T59793	dbEST
Sequence 4578	H19153	dbEST	Sequence 4630	T62753	dbEST
Sequence 4579	H19367	dbEST	Sequence 4631	T65174	dbEST
Sequence 4580	H20888	dbEST	Sequence 4632	T65642	dbEST
Sequence 4581	H23370	dbEST	Sequence 4633	T68057	dbEST
Sequence 4582	H25211	dbEST	Sequence 4634	T71122	dbEST
Sequence 4583	H25825	dbEST	Sequence 4635	T74004	
Sequence 4584	H26319	dbEST	Sequence 4636	T79957	dbEST
Sequence 4585	H28581	dbEST	Sequence 4637	T81666	dbEST
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Sequence 4587	H29635	dbEST	Sequence 4639	W01739	dbEST
Sequence 4588	H39809	dbEST	Sequence 4640	W02575	dbEST
Sequence 4589	H49903	dbEST	Sequence 4641	W04588	dbEST
Sequence 4590	H54240	dbEST	Sequence 4642	W28466	dbEST
Sequence 4591	H60247	dbEST	Sequence 4643	W31167	dbEST
Sequence 4592	H61232	dbEST	Sequence 4644	W32156	dbEST
Sequence 4593	H70543	dbEST	Sequence 4645	W38407	dbEST
Sequence 4594	H71076	dbEST	Sequence 4646		dbEST
Sequence 4595	H79973	dbEST	Sequence 4647	W38913	dbEST
Sequence 4596	H83317	dbEST	Sequence 4648	W39053	dbEST
Sequence 4597	H83328	dbEST	Sequence 4649	W48827 W72729	dbEST
Sequence 4598	H85200	dbEST	Sequence 4650	W77826	dbEST
Sequence 4599	H87807	dbEST	Sequence 4651	W77820 W79340	dbEST
Sequence 4600	H89883	dbEST	Sequence 4652	W80673	dbEST
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Sequence 4602	H99586	dbEST	Sequence 4654	Z33579	dbEST dbEST
Sequence 4603	N21201	dbEST	Sequence 4655	N91259	NUCPATENT
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Sequence 4605	N26421	dbEST	Sequence 4657	Q37811	NUCPATENT
Sequence 4606	N28008	dbEST	Sequence 4658	Q65520	NUCPATENT
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Sequence 4611	N47360	dbEST	Sequence 4663	T79274	NUCPATENT
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Sequence 4613	R11885	dbEST	Sequence 4665	V20443	NUCPATENT
Sequence 4614	R13878	dbEST	Sequence 4666	V29267	NUCPATENT
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Sequence 4616	R18274	dbEST	Sequence 4668	V43612	NUCPATENT
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Sequence 4619	R34553	dbEST	Sequence 4671	V58619	NUCPATENT
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Sequence 4621	R42994	dbEST	Sequence 4673	V59539	NUCPATENT
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Sequence 4623	R59031	dbEST	Sequence 4675	V64158	NUCPATENT
Sequence 4624	R61601	dbEST	Sequence 4676	V69855	NUCPATENT
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Sequence 4677	V70354	NUCPATENT	Sequence 4726	AB002346	GENBANK
Sequence 4678	V72211	NUCPATENT	Sequence 4727	AB002353	GENBANK
Sequence 4679	V81414	NUCPATENT	Sequence 4728	AB002375	GENBANK
Sequence 4680	V84510	NUCPATENT	Sequence 4729	AB002380	GENBANK
Sequence 4681	V86232	NUCPATENT	Sequence 4730	AB002386	GENBANK
Sequence 4682	V89565	NUCPATENT	Sequence 4731	AB007929	GENBANK
Sequence 4683	V90009	NUCPATENT	Sequence 4732	AB007932	GENBANK
Sequence 4684	X00630	NUCPATENT	Sequence 4733	AB007956	GENBANK
Sequence 4685	X03841	NUCPATENT	Sequence 4734	AB011120	GENBANK
Sequence 4686	X07429	NUCPATENT	Sequence 4735	AB011147	GENBANK
Sequence 4687	X20426	NUCPATENT	Sequence 4736	AB014888	GENBANK
Sequence 4688	X20437	NUCPATENT	Sequence 4737	AB016533	GENBANK
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Sequence 4690	X37385	NUCPATENT	Sequence 4739	AB017971	GENBANK
Sequence 4691	X37553	NUCPATENT	Sequence 4740	AB018284	GENBANK
Sequence 4692	X39825	NUCPATENT	Sequence 4741	AB018313	GENBANK
Sequence 4693	X39912	NUCPATENT	Sequence 4742	AB019392	GENBANK
Sequence 4694	X52268	NUCPATENT	Sequence 4743	AB019568	GENBANK
Sequence 4695	X56415	NUCPATENT	Sequence 4744	AB020665	GENBANK
Sequence 4696	X59352	NUCPATENT	Sequence 4745	AB020679	GENBANK
Sequence 4697	X80676	NUCPATENT	Sequence 4746	AB023145	GENBANK
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Sequence 4703	Z15132	NUCPATENT	Sequence 4752	AB029290	GENBANK
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Sequence 4705	Z15987	NUCPATENT	Sequence 4754	AB033033	GENBANK
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Sequence 4709	Z33949	NUCPATENT	Sequence 4758	AF001893	GENBANK
Sequence 4710	AC29436	PREPATNUC	Sequence 4759	AF002210	GENBANK
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	#		Sequence 4775	AF038404	GENBANK
Sequence 4724	A06846	GENBANK	Sequence 4776	AF041483	GENBANK
Sequence 4725	A17546	GENBANK	Sequence 4777	AF045167	GENBANK
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Sequence 4778         AFO47599         GENBANK         Sequence 4831         ALJ049471         GENBANK           Sequence 4780         AF054990         GENBANK         Sequence 4832         AL050035         GENBANK           Sequence 4781         AF064990         GENBANK         Sequence 4833         AL050035         GENBANK           Sequence 4782         AF061255         GENBANK         Sequence 4834         AL080135         GENBANK           Sequence 4783         AF064769         GENBANK         Sequence 4834         AL100273         GENBANK           Sequence 4786         AF070651         GENBANK         Sequence 4838         AL110273         GENBANK           Sequence 4786         AF070651         GENBANK         Sequence 4839         D10040         GENBANK           Sequence 4786         AF070651         GENBANK         Sequence 4839         D1366         GENBANK           Sequence 4787         AF070657         GENBANK         Sequence 4839         D1366         GENBANK           Sequence 4780         AF071010         GENBANK         Sequence 4839         D17409         GENBANK           Sequence 4791         AF074016         GENBANK         Sequence 4841         D17409         GENBANK           Sequence 4793						
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Sequence 4785         AF070523         GENBANK         Sequence 4837         AL117471         GENBANK           Sequence 4786         AF070561         GENBANK         Sequence 4838         D10040         GENBANK           Sequence 4786         AF070657         GENBANK         Sequence 4838         D13866         GENBANK           Sequence 4788         AF070657         GENBANK         Sequence 4840         D16937         GENBANK           Sequence 4789         AF071202         GENBANK         Sequence 4841         D17260         GENBANK           Sequence 4790         AF074016         GENBANK         Sequence 4842         D17409         GENBANK           Sequence 4791         AF077030         GENBANK         Sequence 4842         D17409         GENBANK           Sequence 4792         AF078858         GENBANK         Sequence 4844         D29012         GENBANK           Sequence 4793         AF08455         GENBANK         Sequence 4845         D31763         GENBANK           Sequence 4793         AF085243         GENBANK         Sequence 4845         D33710         GENBANK           Sequence 4793         AF085584         GENBANK         Sequence 4851         D6371         GENBANK           Sequence 4793         A	Sequence 4782	AF061258	GENBANK	•	AL080135	GENBANK
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Sequence 4827 AJ010482 GENBANK Sequence 4879 L38951 GENBANK Sequence 4888 AJ132583 GENBANK Sequence 4880 M10119 GENBANK						
Sequence 4828 AJ132583 GENBANK Sequence 4880 M10119 GENBANK				-		
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Sequence 4829 AJ224172 GENBANK Sequence 4881 M10906 GENBANK	Sequence 4829	AJ224172	GENBANK	Sequence 4881	M10906	GENBANK

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Sequence 4885	M13692	GENBANK	Sequence 4937	U794 <b>57</b>	GENBANK
Sequence 4886	M15395	GENBANK	Sequence 4938	U82226	GENBANK
Sequence 4887	M15885	GENBANK	Sequence 4939	U82256	GENBANK
Sequence 4888	M16247	GENBANK	Sequence 4940	U94586	GENBANK
Sequence 4889	M17885	GENBANK	Sequence 4941	X01037	GENBANK
Sequence 4890	M17886	GENBANK	Sequence 4942	X01630	GENBANK
Sequence 4891	M21895	GENBANK	Sequence 4943	X04098	GENBANK
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Sequence 4893	M22918	GENBANK	Sequence 4945	X04408	GENBANK
Sequence 4894	M26325	GENBANK	Sequence 4946	X04526	GENBANK
Sequence 4895	M26663	GENBANK	Sequence 4947	X13839	GENBANK
Sequence 4896	M26880	GENBANK	Sequence 4948	X15729	GENBANK
Sequence 4897	M29064	GENBANK	Sequence 4949	X57352	GENBANK
Sequence 4898	M33308	GENBANK	Sequence 4950	X58141	GENBANK
Sequence 4899	M34840	GENBANK	Sequence 4951	X60036	GENBANK
Sequence 4900	M36647	GENBANK	Sequence 4952	X63380	GENBANK
Sequence 4901	M60255	GENBANK	Sequence 4953	X63422	GENBANK
Sequence 4902	M61832	GENBANK	Sequence 4954	X63564	GENBANK
Sequence 4903	M64098	GENBANK	Sequence 4955	X68277	GENBANK
Sequence 4904	M64241	GENBANK	Sequence 4956	X69711	GENBANK
Sequence 4905	M64571	GENBANK	Sequence 4957	X73874	GENBANK
Sequence 4906	M68840	GENBANK	Sequence 4958	X76302	GENBANK
Sequence 4907	M81757	GENBANK	Sequence 4959	X76732	GENBANK
Sequence 4908	M94345	GENBANK	Sequence 4960	X81889	GENBANK
Sequence 4909	S50015	GENBANK	Sequence 4961	X84908	GENBANK
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Sequence 4933	U52969	GENBANK	Sequence 4985	AA132342	dbEST

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Sequence 4991	AA178907	dbEST	Sequence 5043	AA359763	dbEST
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Sequence 5037	AA339065	dbEST	Sequence 5089	AA578265	<b>TRAdt</b>

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Sequence 5137	Al246497	dbEST	Sequence 5189	AL042536	dbEST
Sequence 5138	AI253395	dbEST	Sequence 5190	AW003287	dbEST
Sequence 5139	Al267158	dbEST	Sequence 5191	AW024797	dbEST
Sequence 5140	Al267216	dbEST	Sequence 5192	AW071906	dbEST
Sequence 5141	Al267282	dbEST	Sequence 5193	AW130325	dbEST

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		" 507			
Sequence 5194		dbEST	Sequence 5246	Z17710	NUCPATENT
Sequence 5195	AW177929	dbEST	Sequence 5247	Z33671	NUCPATENT
Sequence 5196	AW177931	dbEST			
Sequence 5197	C05816	dbEST	Sequence #	Accession	Database
Sequence 5198	C18484	dbEST	50.40	#	
Sequence 5199	D58694	dbEST	Sequence 5248	A06846	GENBANK
Sequence 5200	F24428	db <b>EST</b>	Sequence 5249	A06919	GENBANK
Sequence 5201	F27796	dbEST	Sequence 5250	A06925	GENBANK
Sequence 5202	H00112	dbEST	Sequence 5251	A21185	GENBANK
Sequence 5203	H08924	dbEST	Sequence 5252	A23013	GENBANK
Sequence 5204	H18732	dbEST	Sequence 5253	AB000220	GENBANK
Sequence 5205	H58534	dbEST	Sequence 5254	AB000584	GENBANK
Sequence 5206	M85357	dbEST	Sequence 5255	AB002312	GENBANK
Sequence 5207	N40017	dbEST	Sequence 5256	AB002350	GENBANK
Sequence 5208	N86292	dbEST	Sequence 5257	AB002382	GENBANK
Sequence 5209	R70270	dbEST	Sequence 5258	AB002806	GENBANK
Sequence 5210	R99478	dbEST	Sequence 5259	AB004851	GENBANK
Sequence 5211	W00471	dbEST	Sequence 5260	AB005535	GENBANK
Sequence 5212	W04380	dbEST	Sequence 5261	AB006534	GENBANK
Sequence 5213	W07165	dbEST	Sequence 5262	AB006621	GENBANK
Sequence 5214	W37945	dbEST	Sequence 5263	AB007867	GENBANK
Sequence 5215	W73588	dbEST	Sequence 5264	AB007882	GENBANK
Sequence 5216	W78967	dbEST	Sequence 5265	AB007892	GENBANK
Sequence 5217	W85764	dbEST	Sequence 5266	AB007931	GENBANK
Sequence 5218	N60172	NUCPATENT	Sequence 5267	AB007947	GENBANK
Sequence 5219	N91825	NUCPATENT	Sequence 5268	AB008226	GENBANK
Sequence 5220	Q43526	NUCPATENT	Sequence 5269	AB011164	<b>GENBAN</b> <sup>™</sup>
Sequence 5221	Q66256	NUCPATENT	Sequence 5270	AB014511	GENBANA
Sequence 5222	Q74061	NUCPATENT	Sequence 5271	AB014542	GENBANK
Sequence 5223	T00880	NUCPATENT	Sequence 5272	AB014552	GENBANK
Sequence 5224	T04864	NUCPATENT	Sequence 5273	AB018274	GENBANK
Sequence 5225	T36781	NUCPATENT	Sequence 5274	AB018284	GENBANK
Sequence 5226	T59998	NUCPATENT	Sequence 5275	AB018301	GENBANK
Sequence 5227	T89060	NUCPATENT	Sequence 5276	AB018315	GENBANK
Sequence 5228	T91054	NUCPATENT	Sequence 5277	AB018319	GENBANK
Sequence 5229	V07571	NUCPATENT	Sequence 5278	AB018340	GENBANK
Sequence 5230	V34212	NUCPATENT	Sequence 5279	AB018352	GENBANK
Sequence 5231	V45442	NUCPATENT	Sequence 5280	AB019568	GENBANK
Sequence 5232	V64828	NUCPATENT	Sequence 5281	AB019691	GENBANK
Sequence 5233	V69855	NUCPATENT	Sequence 5282	AB020637	GENBANK
Sequence 5234	V71075	NUCPATENT	Sequence 5283	AB020652	GENBANK
Sequence 5235	V71742	NUCPATENT	Sequence 5284	AB020692	GENBANK
Sequence 5236	V74071	NUCPATENT	Sequence 5285	AB020698	GENBANK
Sequence 5237	V84395	NUCPATENT	Sequence 5286	AB021288	GENBANK
Sequence 5238	V86225	NUCPATENT	Sequence 5287	AB023150	GENBANK
Sequence 5239	V86232	NUCPATENT	Sequence 5288	AB023153	GENBANK
Sequence 5240	X04125	NUCPATENT	Sequence 5289	AB027466	GENBANK
Sequence 5241	X87624	NUCPATENT	Sequence 5290	AB028964	GENBANK
Sequence 5242	X98561	NUCPATENT	Sequence 5291	AB028971	GENBANK
Sequence 5243	X99098		Sequence 5292	AB029000	GENBANK
Sequence 5244	Z06250	NUCPATENT	Sequence 5293	AB029027	GENBANK
Sequence 5245		NUCPATENT	Sequence 5294	AB032957	GENBANK
Sequence 5245	Z11491	NUCPATENT	Coquence 0234	, 2002001	JE(15) (11)

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Sequence 5295	AB033010	GENBANK	Sequence 5347	AF062318	GENBANK
Sequence 5296	AB033017	GENBANK	Sequence 5348	AF064491	GENBANK
Sequence 5297	AB033070	GENBANK	Sequence 5349	AF064819	GENBANK
Sequence 5298	AB033074	GENBANK	Sequence 5350	AF065388	GENBANK
Sequence 5299	AF007791	GENBANK	Sequence 5351	AF068 <b>754</b>	GENBANK
Sequence 5300	AF008443	GENBANK	Sequence 5352	AF069073	GENBANK
Sequence 5301	AF013759	GENBANK	Sequence 5353	AF069601	GENBANK
Sequence 5302	AF015283	GENBANK	Sequence 5354	AF070523	GENBANK
Sequence 5303	AF017305	GENBANK	Sequence 5355	AF070548	GENBANK
Sequence 5304	AF017445	GENBANK	Sequence 5356	AF070553	GENBANK
Sequence 5305	AF019226	GENBANK	Sequence 5357	AF070554	GENBANK
Sequence 5306	AF020038	GENBANK	Sequence 5358	AF070562	GENBANK
Sequence 5307	AF020591	GENBANK	Sequence 5359	AF070596	GENBANK
Sequence 5308	AF020768	GENBANK	Sequence 5360	AF070652	GENBANK
Sequence 5309	AF021336	GENBANK	Sequence 5361	AF071202	GENBANK
Sequence 5310	AF023676	GENBANK	Sequence 5362	AF072371	GENBANK
	AF025998	GENBANK	Sequence 5363	AF077030	GENBANK
Sequence 5311	AF025336 AF026292	GENBANK	Sequence 5364	AF077037	GENBANK
Sequence 5312	AF020292 AF027302	GENBANK	Sequence 5365	AF077202	GENBANK
Sequence 5313	AF034209	GENBANK	Sequence 5366	AF078854	GENBANK
Sequence 5314	AF034795	GENBANK	Sequence 5367	AF078858	GENBANK
Sequence 5315	AF034795 AF038172	GENBANK	Sequence 5368	AF078863	GENBANK
Sequence 5316			Sequence 5369	AF082858	GENBANK
Sequence 5317	AF038175	GENBANK	•	AF082889	GENBANK
Sequence 5318	AF038404	GENBANK	Sequence 5370	AF083190	GENBANK
Sequence 5319	AF038451	GENBANK	Sequence 5371	AF086168	GENBANK
Sequence 5320	AF038962	GENBANK	Sequence 5372		
Sequence 5321	AF039081	GENBANK	Sequence 5373	AF086245	GENBANK
Sequence 5322	AF039918	GENBANK	Sequence 5374	AF086336	GENBANK
Sequence 5323	AF040250	GENBANK	Sequence 5375	AF087020	GENBANK
Sequence 5324	AF040958	GENBANK	Sequence 5376	AF089747	GENBANK
Sequence 5325	AF042166	GENBANK	Sequence 5377	AF091084	GENBANK
Sequence 5326	AF042284	GENBANK	Sequence 5378	AF092135	GENBANK
Sequence 5327	AF042384	GENBANK	Sequence 5379	AF096773	GENBANK
Sequence 5328	AF045583	GENBANK	Sequence 5380	AF096774	GENBANK
Sequence 5329	AF047020	GENBANK	Sequence 5381	AF097514	GENBANK
Sequence 5330	AF047185	GENBANK	Sequence 5382	AF099149	GENBANK
Sequence 5331	AF047436	GENBANK	Sequence 5383	AF100741	GENBANK
Sequence 5332	AF048731	GENBANK	Sequence 5384	AF100756	GENBANK GENBANK
Sequence 5333	AF050641	GENBANK	Sequence 5385	AF102846	
Sequence 5334	AF051321	GENBANK	Sequence 5386	AF103907	GENBANK
Sequence 5335	AF052093	GENBANK	Sequence 5387	AF112227	GENBANK
Sequence 5336	AF052149	GENBANK	Sequence 5388	AF126181	GENBANK GENBANK
Sequence 5337	AF052175	GENBANK	Sequence 5389	AF128528	
Sequence 5338	AF053070	GENBANK	Sequence 5390	AF131220	GENBANK
Sequence 5339	AF054175	GENBANK	Sequence 5391	AF132940	GENBANK
Sequence 5340	AF054181	GENBANK	Sequence 5392	AF138300	GENBANK
Sequence 5341	AF054838	GENBANK	Sequence 5393	AF147331	GENBANK
Sequence 5342	AF054988	GENBANK	Sequence 5394	AF147334	GENBANK
Sequence 5343	AF057706	GENBANK	Sequence 5395	AF149045	GENBANK
Sequence 5344	AF058718	GENBANK	Sequence 5396	AF151105	GENBANK
Sequence 5345	AF060219	GENBANK	Sequence 5397	AF151800	GENBANK
Sequence 5346	AF061258	GENBANK	Sequence 5398	AF151840	GENBANK

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			_		
Sequence 5399	AF151846	GENBA <b>NK</b>	Sequence 5451	D28482	GENBANK
Sequence 5400	AF1518 <b>61</b>	GENBANK	Sequence 5452	<b>D2</b> 9640	GENBANK
Sequence 5401	AF1518 <b>68</b>	GENBANK	Sequence 5453	D29954	GENBANK
Sequence 5402	AF151884	GENBANK	Sequence 5454	<b>D3</b> 1886	GENBANK
Sequence 5403	AF151893	GENBANK	Sequence 5455	D38037	GENBANK
Sequence 5404	AF159056	GENBANK	Sequence 5456	D38549	GENBANK
Sequence 5405	AF159092	GENBANK	Sequence 5457	D42039	GENBANK
Sequence 5406	AF159295	GENBANK	Sequence 5458	D42044	GENBANK
Sequence 5407	AF168956	GENBANK	Sequence 5459	D42047	GENBANK
Sequence 5408	AF177291	GENBANK	Sequence 5460	D42138	GENBANK
Sequence 5409	AF177775	GENBANK	Sequence 5461	D43950	GENBANK
Sequence 5410	AF188298	GENBANK	Sequence 5462	D49372	GENBANK
Sequence 5411	AF188745	GENBANK	Sequence 5463	D49489	GENBANK
Sequence 5412	AF188746	GENBANK	Sequence 5464	D50369	GENBANK
Sequence 5413	AF190167	GENBANK	Sequence 5465	D50371	GENBANK
Sequence 5414	AJ010046	GENBANK	Sequence 5466	D50579	GENBANK
Sequence 5415	AJ010482	GENBANK	Sequence 5467	D50916	GENBANK
Sequence 5416	AJ012463	GENBANK	Sequence 5468	D50919	GENBANK
Sequence 5417	AJ012499	GENBANK	Sequence 5469	D55653	GENBANK
Sequence 5418	AJ130733	GENBANK	Sequence 5470	D63486	GENBANK
Sequence 5419	AJ223075	GENBANK	Sequence 5471	D64110	GENBANK
Sequence 5420	AJ223350	GENBANK	Sequence 5472	D80009	GENBANK
Sequence 5421	AJ224172	GENBANK	Sequence 5473	D82060	GENBANK
Sequence 5422	AJ249731	GENBANK	Sequence 5474	D83032	GENBANK
Sequence 5423	AL049705	GENBANK	Sequence 5475	D83735	GENBANK
Sequence 5424	AL049969	GENBANK	Sequence 5476	D85181	GENBANK
Sequence 5425	AL050198	GENBANK	Sequence 5477	D86984	GENBANK
Sequence 5426	AL050274	GENBANK	Sequence 5478	D87127	GENBANK
Sequence 5427	AL080089	GENBANK	Sequence 5479	D87292	GENBANK
Sequence 5428	AL080135	GENBANK	Sequence 5480	D87453	GENBANK
Sequence 5429	AL080209	GENBANK	Sequence 5481	D87462	GENBANK
Sequence 5430	AL096880	GENBANK	Sequence 5482	D87666	GENBANK
Sequence 5431	AL109672	GENBANK	Sequence 5483	D87684	GENBANK
Sequence 5432	AL110183	GENBANK	Sequence 5484	D87735	GENBANK
Sequence 5433	AL117413	GENBANK	Sequence 5485	D87969	GENBANK
Sequence 5434	AL117550	GENBANK	Sequence 5486	D88674	GENBANK
Sequence 5435	AL117595	GENBANK	Sequence 5487	D90373	GENBANK
Sequence 5436	AL121733	GENBANK	Sequence 5488	D90427	GENBANK
Sequence 5437	D00422	GENBANK	Sequence 5489	E01650	GENBANK
Sequence 5438	D13641	GENBANK	Sequence 5490	E01915	GENBANK
Sequence 5439	D13866	GENBANK	Sequence 5491	E02628	GENBANK
Sequence 5440	D13900	GENBANK	Sequence 5492	E03413	GENBANK
Sequence 5441	D14662	GENBANK	Sequence 5493	E06721	GENBANK
Sequence 5442	D14664	GENBA <b>NK</b>	Sequence 5494	J00194	GENBANK
Sequence 5443	D16111	GENBANK	Sequence 5495	J02642	GENBANK
Sequence 5444	D16922	GENBANK	Sequence 5496	J02876	GENBANK
Sequence 5445	D17409	GENBANK	Sequence 5497	J02908	GENBANK
Sequence 5446	D17554	GENBANK	Sequence 5498	J03007	GENBANK
Sequence 5447	D21243	GENBANK	Sequence 5499	J03015	GENBANK
Sequence 5448	D21851	GENBANK	Sequence 5500	J03171	GENBANK
Sequence 5449	D21853	GENBANK	Sequence 5501	J03248	GENBANK
Sequence 5450	D26068	GENBANK	Sequence 5502	J03507	GENBANK
_ = = = = = = = = = = = = = = = = = = =	525000	C			

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				_		
Sequence 5503	J03592	GENBANK	Sequ	uence 555 <b>5</b>	M26663	GENBANK
Sequence 5504	J03799	GENBA <b>NK</b>	Sequ	uence 5556	M27110	GENBANK
Sequence 5505	J03870	GENBA <b>NK</b>	Sequ	uence 5557	M28016	GENBANK
Sequence 5506	J05633	GENBA <b>N</b> K	Sequ	uence 55 <b>58</b>	M28211	GENBANK
Sequence 5507	K00409	GENBANK	Seq	uence 55 <b>59</b>	M29366	GENBANK
Sequence 5508	K01911	GENBANK	Seq	uence 55 <b>60</b>	M29872	GENBANK
Sequence 5509	L05092	GENBA <b>NK</b>	Seq	uence 5561	M31212	GENBANK
Sequence 5510	L05093	GENBANK	Seq	uence 55 <b>62</b>	M31606	GENBANK
Sequence 5511	L05779	GENBANK	Seq	uence 55 <b>63</b>	M33146	GENBANK
Sequence 5512	L07033	GENBANK	Seq	uence 5564	M33308	GENBANK
Sequence 5513	L07517	GENBANK	Seq	uence 55 <b>65</b>	M34840	GENBANK
Sequence 5514	L08044	GENBANK	Seq	uence 5566	M36634	GENBANK
Sequence 5515	L12136	GENBANK	Seq	uence 5567	M57399	GENBANK
Sequence 5516	L15203	GENBANK	Seq	uence 5568	M57567	GENBANK
Sequence 5517	L16785	GENBANK	Seq	uence 5569	M58028	GENBANK
Sequence 5518	L19597	GENBANK	Seq	uence 5570	M58485	GENBANK
Sequence 5519	L19605	GENBANK		uence 5571	M58549	GENBANK
Sequence 5520	L19956	GENBANK	Seq	uence 5572	M60457	GENBANK
Sequence 5521	L20773	GENBANK	Seq	uence 5573	M60857	GENBANK
Sequence 5522	L20941	GENBANK	Seq	uence 5574	M61832	GENBANK
Sequence 5523	L23320	GENBANK	Seq	uence 5575	M63573	GENBANK
Sequence 5524	L28010	GENBANK	Seq	uence 5576	M64098	GENBANK
Sequence 5525	L29008	GENBANK	Seq	uence 5577	M64241	GENBANK
Sequence 5526	L31951	GENBANK	Seq	uence 5578	M64571	GENBANK
Sequence 5527	L34840	GENBANK	Seq	uence 5579	M69066	GENBANK
Sequence 5528	L37080	GENBANK	Sed	uence 5580	M69180	GENBANK
Sequence 5529	L38486	GENBANK	Sed	uence 5581	M74777	GENBANK
Sequence 5530	L38961	GENBANK	Sed	uence 5582	M76180	GENBANK
Sequence 5531	L38995	GENBANK	Sed	uence 5583	M77830	GENBANK
Sequence 5532	L42531	GENBANK	Sec	uence 5584	M81757	GENBANK
Sequence 5533	L47665	GENBANK	Sec	quence 5585	M86752	GENBANK
Sequence 5534	M10036	GENBANK		quence 55 <b>86</b>	M88279	GENBANK
Sequence 5535	M10119	GENBANK		риепсе 5587	M92381	GENBANK
Sequence 5536	M13520	GENBANK		quence 5588	M92843	GENBANK
Sequence 5537	M14335	GENBANK	Sec	quence 5589	M95178	GENBANK
Sequence 5538	M15470	GENBANK		quence 5590	M99701	GENBANK
Sequence 5539	M15887	GENBANK		quence 5591	S60099	GENBANK
Sequence 5540	M16660	GENBANK		quence 5592	S67325	GENBANK
Sequence 5541	M16804	GENBANK		quence 5593	S69022	GENBANK
Sequence 5542	M17851	GENBANK	Sed	quence 5594	S69272	GENBANK
Sequence 5543	M19308	GENBANK		quence 5595	S69738	GENBANK
Sequence 5544	M20471	GENBANK		quence 5596	S70154	GENBANK
Sequence 5545	M21895	GENBANK		quence 5597	S73591	GENBANK
Sequence 5546	M21896	GENBANK		quence 5598	S74678	GENBANK
Sequence 5547	M22324	GENBANK	Sec	quence 5599	S75755	GENBANK
Sequence 5548	M24194	GENBANK		quence 5600	U02390	GENBANK
Sequence 5549	M24795	GENBANK		quence 5601	U04209	GENBANK
Sequence 5550	M24902	GENBANK		quence 5602	U05291	GENBANK
Sequence 5551	M24906	GENBANK		quence 5603	U06632	GENBANK
Sequence 5552	M25246	GENBANK		quence 5604	U07919	GENBANK
Sequence 5553	M26324	GENBANK		quence 5605	U09550	GENBANK
Sequence 5554	M26325	GENBANK	Se	quence 5606	U14510	GENBANK
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Sequence 5607	U14750	GENBANK	Sequence 5659	X53777	GENBANK
Sequence 5608	U14968	GENBANK	Sequence 5660	X56998	GENBANK
Sequence 5609	U14969	GENBANK	Sequence 5661	X57398	GENBANK
Sequence 5610	U14970	GENBANK	Sequence 5662	X60221	GENBANK
Sequence 5611	U16850	GENBANK	Sequence 5663	X60708	GENBANK
Sequence 5612	U16997	GENBANK	Sequence 5664	X66276	GENBANK
Sequence 5613	U17743	GENBANK	Sequence 5665	X66534	GENBANK
Sequence 5614	U18062	GENBANK	Sequence 5666	X67951	GENBANK
Sequence 5615	U18197	GENBANK	Sequence 5667	X70394	GENBANK
Sequence 5616	U18291	GENBANK	Sequence 5668	X71129	GENBANK
Sequence 5617	U18543	GENBANK	Sequence 5669	X73114	GENBANK
Sequence 5618	U27143	GENBANK	Sequence 5670	X74070	GENBANK
Sequence 5619	U28424	GENBANK	Sequence 5671	X75546	GENBANK
Sequence 5620	U30313	GENBANK	Sequence 5672	X75593	GENBANK
Sequence 5621	U30826	GENBANK	Sequence 5673	X75861	GENBANK
Sequence 5622	U31382	GENBANK	Sequence 5674	X76105	GENBANK
Sequence 5623	U31905	GENBANK	Sequence 5675	X76538	GENBANK
Sequence 5624	U36764	GENBANK	Sequence 5676	X77196	GENBANK
Sequence 5625	U37519	GENBANK	Sequence 5677	X78137	GENBANK
Sequence 5626	U39360	GENBANK	Sequence 5678	X78520	GENBANK
Sequence 5627	U39400	GENBANK	Sequence 5679	X79389	GENBANK
Sequence 5628	U40272	GENBANK	Sequence 5680	X79888	GENBANK
Sequence 5629	U43701	GENBANK	Sequence 5681	X80822	GENBANK
Sequence 5630	U44839	GENBANK	Sequence 5682	X82157	GENBANK
Sequence 5631	U46751	GENBANK	Sequence 5683	X85373	GENBANK
Sequence 5632	U51903	GENBANK	Sequence 5684	X87241	GENBANK
Sequence 5633	U54562	GENBANK	Sequence 5685	X90583	GENBANK
Sequence 5634	U54996	GENBANK	Sequence 5686	X92098	GENBANK
Sequence 5635	U58855	GENBANK	Sequence 5687	X93036	GENBANK
Sequence 5636	U66615	GENBANK	Sequence 5688	X97065	GENBANK
Sequence 5637	U67171	GENBANK	Sequence 5689	X97674	GENBANK
Sequence 5638	U69263	GENBANK	Sequence 5690	X98801	GENBANK
Sequence 5639	U69668	GENBANK	Sequence 5691	Y00282	GENBANK
Sequence 5640	U79241	GENBANK	Sequence 5692	Y07968	GENBANK
Sequence 5641	U79304	GENBANK	Sequence 5693	Y12781	GENBANK
Sequence 5642	U80456	GENBANK	Sequence 5694	Y13936	GENBANK
Sequence 5643	U83460	GENBANK	Sequence 5695	Y16132	GENBANK
Sequence 5644	U95646	GENBANK	Sequence 5696	Y16610	GENBANK
Sequence 5645	U96721	GENBANK	Sequence 5697	Z26317	GENBANK
Sequence 5646	X00351	GENBANK	Sequence 5698	Z46973	GENBANK
Sequence 5647	X04106	GENBANK	Sequence 5699	Z50749	GENBANK
Sequence 5648	X04325	GENBANK	Sequence 5700	AA001299	dbEST
Sequence 5649	X04408	GENBANK	Sequence 5701	AA001316	dbEST
Sequence 5650	X04571	GENBANK	Sequence 5702	AA002190	dbEST
Sequence 5651	X05332	GENBANK	Sequence 5703	AA009641	dbEST
Sequence 5652	X06617	GENBANK	Sequence 5704	AA010236	dbEST
Sequence 5653	X07876	GENBANK	Sequence 5705	AA010607	dbEST
Sequence 5654	X13839	GENBANK	Sequence 5706	AA013094	dbEST
Sequence 5655	X14608	GENBANK	Sequence 5707	AA015605	dbEST
Sequence 5656	X15187	GENBANK	Sequence 5708	AA018347	dbEST
Sequence 5657	X16940	GENBANK	Sequence 5709	AA019377	dbEST
Sequence 5658	X17206	GENBANK	Sequence 5710	AA022583	dbEST
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					JE COT
Sequence 5711	AA024398	dbEST	Sequence 5763	AA074683	dbEST
Sequence 5712	AA024532	dbEST	Sequence 5764	AA074819	dbEST
Sequence 5713	AA024967	dbEST	Sequence 5765	AA075474	dbEST
Sequence 5714	AA025731	dbEST	Sequence 5766	AA075600	dbEST
Sequence 5715	AA025875	dbEST	Sequence 5767	AA075781	dbEST
Sequence 5716	AA0:26658	dbEST	Sequence 5768	AA075922	dbEST
Sequence 5717	AA029143	dbEST	Sequence 5769	AA078218	dbEST
Sequence 5718	AA029157	dbEST	Sequence 5770	AA079346	dbEST
Sequence 5719	AA029584	dbEST	Sequence 5771	AA079524	dbEST
Sequence 5720	AA033517	dbEST	Sequence 5772	AA079690	dbEST
Sequence 5721	AA034255	dbEST	Sequence 5773	AA079755	dbEST
Sequence 5722	AA034946	dbEST	Sequence 5774	AA081708	dbEST
Sequence 5723	AA035773	dbEST	Sequence 5775	AA081812	dbEST
Sequence 5724	AA037217	dbEST	Sequence 5776	AA081842	dbEST
Sequence 5725	AA037857	dbEST	Sequence 5777	AA082839	dbEST
Sequence 5726	AA039546	dbEST	Sequence 5778	AA083008	dbEST
Sequence 5727	AA040140	dbEST	Sequence 5779	AA083191	dbEST
Sequence 5728	AA044004	dbEST	Sequence 5780	AA083345	dbEST
Sequence 5729	AA044291	dbEST	Sequence 5781	AA083410	dbEST
Sequence 5730	AA044570	dbEST	Sequence 5702	AA083603	dbEST
Seguence 5731	AA045185	dbEST	Sequence 5783	AA08 <b>4068</b>	dbEST
Sequence 5732	AA045342	dbEST	Sequence 5784	AA084552	dbEST
Sequence 5733	AA045388	dbEST	Sequence 5785	AA084560	dbEST
Sequence 5734	AA045741	dbEST	Sequence 5786	AA086326	dbEST
Sequence 5735	AA046040	dbEST	Sequence 5787	AA086377	dbEST
Sequence 5736	AA046483	dbEST	Sequence 5788	AA088544	dbEST
Sequence 5737	AA046836	dbEST	Sequence 5789	AA088603	dbEST
Sequence 5738	AA046973	dbEST	Sequence 5790	AA096453	dbEST
Sequence 5739	AA047015	dbEST	Sequence 5791	AA099875	dbEST
Sequence 5740	AA047304	dbEST	Sequence 5792	AA099917	dbEST
Sequence 5741	AA053250	dbEST	Sequence 5793	AA100195	cbEST
Sequence 5742	AA053399	dbEST	Sequence 5794	AA100852	dbEST
Sequence 5743	AA053475	dbEST	Sequence 5795	AA101799	dbEST
Sequence 5744	AA053514	dbEST	Sequence 5796	AA102280	dbEST
Sequence 5745	AA053557	dbEST	Sequence 5797	AA112936	cbEST
Sequence 5746	AA053747	dbEST	Sequence 5798	AA113403	dbEST
Sequence 5747	AA053757	dbEST	Sequence 5799	AA114014	cbEST
Sequence 5748	AA056462	dbEST	Sequence 5800	AA115512	dbEST
Sequence 5749	AA057029	dbEST	Sequence 5801	AA115838	cbEST
Sequence 5750	AA057243	dbEST	Sequence 5802	AA121072	dbEST
Sequence 5751	AA058605	dbEST	Sequence 5803	AA126009	dbEST
Sequence 5752	AA058899	dbEST	Sequence 5804	AA126750	dbEST
Sequence 5753	AA059309	dbEST	Sequence 5805	AA127135	dbEST
Sequence 5754	AA059355	dbEST	Sequence 5806	AA128104	dbEST
Sequence 5755	AA062994	dbEST	Sequence 5807	AA128324	dbEST
Sequence 5756	AA065036	dbEST	Sequence 5808	AA128498	dbEST
•	AA065214	dbEST	Sequence 5809	AA129245	dbEST
Sequence 5757	AA069457		Sequence 5810	AA129726	dbEST
Sequence 5758	AA069814	dbEST dbEST	Sequence 5811	AA129907	dbEST
Sequence 5759			Sequence 5812	AA129962	dbEST
Sequence 5760	AA071084 AA071439	dbEST	Sequence 5813	AA130229	dbEST
Sequence 5761	AA071439 AA074575	dbEST	Sequence 5814	AA130797	dbEST
Sequence 5762	AA014313	dbEST	Dequence 5014	, , , , , , , , , , , , , , , , , , , ,	

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Sequence 5815	AA130946	dbEST	Sequence 5867	AA179442	dbEST
Sequence 5816	AA131828	dbEST	Sequence 5868	AA181113	dbEST
Sequence 5817	AA132596	dbEST	Sequence 5869	AA181392	dbEST
Sequence 5818	AA132844	dbEST	Sequence 5870	AA182617	dbEST
Sequence 5819	AA133579	dbEST	Sequence 5871	AA186406	dbEST
Sequence 5820	AA134203	dbEST	Sequence 5872	AA186432	dbEST
Sequence 5821	AA134234	dbEST	Sequence 5873	AA186477	dbEST
Sequence 5822	AA134715	dbEST	Sequence 5874	AA187395	dbEST
Sequence 5823	AA135456	dbEST	Sequence 5875	AA187762	dbEST
Sequence 5824	AA135663	dbEST	Sequence 5876	AA188633	dbEST
Sequence 5825	AA136096	dbEST	Sequence 5877	AA190615	dbEST
Sequence 5826	AA137170	dbEST	Sequence 5878	AA190873	dbEST
Sequence 5827	AA143286	dbEST	Sequence 5879	AA190890	dbEST
Sequence 5828	AA143579	dbEST	Sequence 5880	AA191119	dbEST
Sequence 5829	AA146616	dbEST	Sequence 5881	AA191211	dbEST
Sequence 5830	AA146701	dbEST	Sequence 5882	AA192597	dbEST
Sequence 5831	AA147403	dbEST	Sequence 5883	AA192759	dbEST
Sequence 5832	AA147826	dbEST	Sequence 5884	AA193297	dbEST
Sequence 5833	AA148019	dbEST	Sequence 5885	AA193661	dbEST
Sequence 5834	AA148219	dbEST	Sequence 5886	AA194473	dbEST
Sequence 5835	AA149936	dbEST	Sequence 5887	AA195015	dbEST
Sequence 5836	AA150320	dbEST	Sequence 5888	AA195617	dbEST
Sequence 5837	AA150383	dbEST	Sequence 5889	AA195947	dbEST
Sequence 5838	AA150860	dbEST	Sequence 5890	AA196357	dbEST
Sequence 5839	AA151350	dbEST .	Sequence 5891	AA203182	dbEST
Sequence 5840	AA151506	dbEST	Sequence 5892	AA205412	dbEST
Sequence 5841	AA151659	dbEST	Sequence 5893	AA205546	dbEST
Sequence 5842	AA152009	dbEST	Sequence 5894	AA206115	dbEST
Sequence 5843	AA152264	db <b>ES</b> T	Sequence 5895	AA206581	cbEST
Sequence 5844	AA155856	dbEST	Sequence 5896	AA211799	dbEST
Sequence 5845	AA156215	dbEST	Sequence 5897	AA214648	dbEST
Sequence 5846	AA157723	dbEST	Sequence 5898	AA216409	dbEST
Sequence 5847	AA159272	dbEST	Sequence 5899	AA216669	dbEST
Sequence 5848	AA159748	dbEST	Sequence 5900	AA218606	dbEST
Sequence 5849	AA160053	dbEST	Sequence 5901	AA219143	dbEST
Sequence 5850	AA160999	dbEST	Sequence 5902	AA223148	dbEST
Sequence 5851	AA164607	db <b>EST</b>	Sequence 5903	AA224124	dbEST
Sequence 5852	AA164977	dbEST	Sequence 5904	AA224876	dbEST
Sequence 5853	AA165197	dbEST	Sequence 5905	AA224985	dbEST
Sequence 5854	AA166675	dbEST	Sequence 5906	AA225857	dbEST
Sequence 5855	AA167044	dbEST	Sequence 5907	AA226359	dbEST
Sequence 5856	AA167804	dbEST	Sequence 5908	AA226898	dbEST
Sequence 5857	AA171510	dbEST	Sequence 5909	AA227326	dbEST
Sequence 5858	AA171587	dbEST	Sequence 5910	AA227579	dbEST
Sequence 5859	AA172103	dbEST	Sequence 5911	AA227837	dbEST
Sequence 5860	AA172156	dbEST	Sequence 5912	AA228273	dbEST
Sequence 5861	AA173098	dbEST	Sequence 5913	AA228282	dbEST
Sequence 5862	AA173459	dbEST	Sequence 5914	AA228836	dbEST
Sequence 5863	AA173952	dbEST	Sequence 5915	AA228940	dbEST
Sequence 5864	AA176747	dbEST	Sequence 5916	AA228953	dbEST
Sequence 5865	AA176784	dbEST	Sequence 5917	AA229057	dbEST
Sequence 5866	AA176979	dbEST	Sequence 5918	AA229145	dbEST

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Sequence 5919	AA230067	dbEST	Sequence 5971	AA301205	dbEST
Sequence 5920	AA233733	dbEST	Sequence 5972	AA301513	dbEST
Sequence 5921	AA235224	dbEST	Sequence 5973	AA301629	dbEST
Sequence 5922	AA235581	dbEST	Sequence 5974	AA302313	dbEST
Sequence 5923	AA236371	dbEST	Sequence 5975	AA302919	dbEST
Sequence 5924	AA236418	dbEST	Sequence 5976	AA303594	dbEST
Sequence 5925	AA236774	dbEST	Sequence 5977	AA303960	dbEST
Sequence 5926	AA244003	dbEST	Sequence 5978	AA304827	dbEST
Sequence 5927	AA244018	dbEST	Sequence 5979	AA305081	dbEST
Sequence 5928	AA244158	dbEST	Sequence 5980	AA305121	dbEST
Sequence 5929	AA244184	dbEST	Sequence 5981	AA305143	dbEST
Sequence 5930	AA248427	dbEST	Sequence 5982	AA305331	dbEST
Sequence 5931	AA251114	dbEST	Sequence 5983	AA305333	dbEST
Sequence 5932	AA251678	dbEST	Sequence 5984	AA305409	dbEST
Sequence 5933	AA256422	dbEST	Sequence 5985	AA305774	dbEST
Sequence 5934	AA257014	dbEST	Sequence 5986	AA305819	dbEST
Sequence 5935	AA258089	dbEST	Sequence 5987	AA305824	dbEST
Sequence 5936	AA278642	dbEST	Sequence 5988	AA306040	dbEST
Sequence 5937	AA279853	dbEST	Sequence 5989	AA306182	dbEST
Sequence 5938	AA280091	dbEST	Sequence 5990	AA306223	dbEST
Sequence 5939	AA281637	dbEST	Sequence 5991	AA306983	dbEST
Sequence 5940	AA282252	dbEST	Sequence 5992	AA307033	dbEST
Sequence 5941	AA282382	dbEST	Sequence 5993	AA307224	dbEST
Sequence 5942	AA283169	dbEST	Sequence 5994	AA307612	dbEST
Sequence 5943	AA283616	dbEST	Sequence 5995	AA307669	dbEST
Sequence 5944	AA284615	dbEST	Sequence 5996	AA307728	dbEST
Sequence 5945	AA284748	dbEST	Sequence 5997	AA308091	dbEST
Sequence 5946	AA284856	dbEST	Sequence 5998	AA308207	dbEST
Sequence 5947	AA285290	dbEST	Sequence 5999	AA308237	dbEST
Sequence 5948	AA287199	dbEST	Sequence 6000	AA308574	dbEST
Sequence 5949	AA287665	dbEST	Sequence 6001	AA308812	dbEST
Sequence 5950	AA290783	dbEST	Sequence 6002	AA309832	dbEST
Sequence 5951	AA291407	dbEST	Sequence 6003	AA310978	dbEST
Sequence 5952	AA292443	dbEST	Sequence 6004	AA311044	dbEST
Sequence 5953	AA292786	dbEST	Sequ <b>ence 6005</b>	AA311048	dbEST
Sequence 5954	AA293855	dbEST	Sequence 6006	AA311227	dbEST
Sequence 5955	AA295706	dbEST	Sequence 6007	AA311677	dbEST
Sequence 5956	AA295762	dbEST	Sequence 6008	AA311896	dbEST
Sequence 5957	AA296074	dbEST	Sequence 6009	AA312406	dbEST
Sequence 5958	AA296282	dbEST	Sequence 6010	AA312864	dbEST
Sequence 5959	AA297097	dbEST	Sequence 6011	AA313209	dbEST
Sequence 5960	AA297506	dbEST	Sequence 6012	AA313468	dbEST
Sequence 5961	AA297592	dbEST	Sequence 6013	AA313653	dbEST
Sequence 5962	AA297750	dbEST	Sequence 6014	AA314146	dbEST
Sequence 5963	AA298490	dbEST	Sequence 6015	- AA314473	dbEST
Sequence 5964	AA298967	dbEST	Sequence 6016	AA314673	dbEST
Sequence 5965	AA299668	dbEST	Sequence 6017	AA314961	dbEST
Sequence 5966	AA299840	dbEST	Sequence 6018	AA315049	dbEST
Sequence 5967	AA299892	dbEST	Sequence 6019	AA315188	dbEST
Sequence 5968	AA299961	dbEST	Sequence 6020	AA315441	dbEST
Sequence 5969	AA300008	dbEST	Sequence 6021	AA315510	dbEST
Sequence 5970	AA300659	dbEST	Sequence 6022	AA315950	dbEST

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Sequence 6023	AA315968	dbEST	Sequence 6075	AA355308	dbEST
Sequence 6024	AA316059	dbEST	Sequence 6076	AA3564 <b>36</b>	dbEST
Sequence 6025	AA316217	dbEST	Sequence 6077	AA357077	dbEST
Sequence 6026	AA316423	dbEST	Sequence 6078	AA363019	dbEST
Sequence 6027	AA316721	dbEST	Sequence 6079	AA363481	dbEST
Sequence 6028	AA316723	dbEST	Sequence 6080	AA364707	dbEST
Sequence 6029	AA317088	dbEST	Sequence 6081	AA365150	dbEST
Sequence 6030	AA317310	dbEST	Sequence 6082	AA369660	dbEST
Sequence 6031	AA317477	dbEST	Sequence 6083	AA369804	dbEST
Sequence 6032	AA317497	dbEST	Sequence 6084	AA370548	dbEST
Sequence 6033	AA317659	dbEST	Sequence 6085	AA371017	dbEST
Sequence 6034	AA317855	dbEST	Sequence 6086	AA371613	dbEST
Sequence 6035	AA317942	dbEST	Sequence 6087	AA372171	dbEST
Sequence 6036	AA318260	dbEST	Sequence 6088	AA373314	dbEST
Sequence 6037	AA318272	dbEST	Sequence 6089	AA374073	dbEST
Sequence 6038	AA320004	dbEST	Sequence 6090	AA374376	dbEST
Sequence 6039	AA320114	dbEST	Sequence 6091	AA374561	dbEST
Sequence 6040	AA320723	dbEST	Sequence 6092	AA375454	dbEST
Sequence 6041	AA321706	dbEST	Sequence 6093	AA376123	dbEST
Sequence 6042	AA322115	dbEST	Sequence 6094	AA377218	dbEST
Sequence 6043	AA322352	dbEST	Sequence 6095	AA384272	dbEST
Sequence 6044	AA323143	dbEST	Sequence 6096	AA393061	dbEST
Sequence 6045	AA326251	dbEST	Sequence 6097	AA393485	dbEST
Sequence 6046	AA326895	dbEST	Sequence 6098	AA393906	dbEST
Sequence 6047	AA327011	dbEST	Sequence 6099	AA394311	dbEST
Sequence 6048	AA327546	dbEST	Sequence 6100	AA397452	dbEST
Sequence 6049	AA328770	dbEST	Sequence 6101	AA397702	dbEST
Sequence 6050	AA330883	dbEST	Sequence 6102	AA398929	dbEST
Sequence 6051	AA332202	dbEST	Sequence 6103	AA399024	dbEST
Sequence 6052	AA332737	dbEST	Sequence 6104	AA399070	dbEST
Sequence 6053	AA334511	dbEST	Sequence 6105	AA399281	dbEST
Sequence 6054	AA335356	dbEST	Sequence 6106	AA399632	dbEST
Sequence 6055	AA336197	dbEST	Sequence 6107	AA400527	dbEST
Sequence 6056	AA337836	dbEST	Sequence 6108	AA400692	dbEST
Sequence 6057	AA338836	dbEST	Sequence 6109	AA401501	dbEST
Sequence 6058	AA339065	dbEST	Sequence 6110	AA401520	dbEST
Sequence 6059	AA340095	dbEST	Sequence 6111	AA401809	dbEST
Sequence 6060	AA340608	dbEST	Sequence 6112	AA404376	dbEST
Sequence 6061	AA340634	dbEST	Sequence 6113	AA404722	dbEST
Sequence 6062	AA340927	dbEST	Sequence 6114	AA411585	dbEST
Sequence 6063	AA341690	dbEST	Sequence 6115	AA412302	dbEST
Sequence 6064	AA341987	dbEST	Sequence 6116	AA412500	dbEST
Sequence 6065	AA343532	dbEST	Sequence 6117	AA417643	dbEST
Sequence 6066	AA344600	dbEST	Sequence 6118	AA417842	dbEST
Sequence 6067	AA345762	dbEST	Sequence 6119	AA420435	dbEST
Sequence 6068	AA345876	dbEST	Sequence 6120	AA420462	dbEST
Sequence 6069	AA346253	dbEST	Sequence 6121	AA420758	dbEST
Sequence 6070	AA347236	dbEST	Sequence 6122	AA420800	dbEST
Sequence 6071	AA347436	dbEST	Sequence 6123	AA420825	dbEST
Sequence 6071	AA352062		Sequence 6124	AA420845	dbEST
Sequence 6073	AA354376	dbEST	•	AA422173	dbEST
•	AA355036	dbEST	Sequence 6125	AA424243	dbEST
Sequence 6074	~~333030	dbEST	Sequence 6126	MM424243	UDES I

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Sequence 6127	AA425212	dbEST	Sequence 6179	AA469348	dbEST
Sequence 6128	AA425305	dbEST	Sequence 6180	AA469926	dbEST
Sequence 6129	AA425460	dbEST	Sequence 6181	AA469979	dbEST
Sequence 6130	AA425492	dbEST	Sequence 6182	AA470575	dbEST
Sequence 6131	AA425509	dbEST	Sequence 6183	AA470602	dbEST
Sequence 6132	AA425583	dbEST	Sequence 6184	AA471056	dbEST
Sequence 6133	AA426351	dbEST	Sequence 6185	AA477075	dbEST
Sequence 6134	AA428216	dbEST	Sequence 6186	AA477397	dbEST
Sequence 6135	AA431579	dbEST	Sequence 6187	AA477803	dbEST
Sequence 6136	AA433869	dbEST	Sequence 6188	AA478415	dbEST
Sequence 6137	AA434286	dbEST	Sequence 6189	AA479151	dbEST
Sequence 6138	AA435894	dbEST	Sequence 6190	AA479167	dbEST
Sequence 6139	AA436394	dbEST	Sequence 6191	AA480124	dbEST
Sequence 6140	AA436652	dbEST	Sequence 6192	AA480282	dbEST
Sequence 6141	AA437304	dbEST	Sequence 6193	AA480529	dbEST
Sequence 6142	AA442017	dbEST	Sequence 6194	AA481738	dbEST
Sequence 6143	AA442145	dbEST	Sequence 6195	AA484711	dbEST
Sequence 6144	AA442287	dbEST	Sequence 6196	AA484821	dbEST
Sequence 6145	AA442517	dbEST	Sequence 6197	AA485467	dbEST
Sequence 6146	AA443757	dbEST	Sequence 6198	AA485853	dbEST
Sequence 6147	AA443834	dbEST	Sequence 6199	AA486619	dbEST
Sequence 6148	AA444072	dbEST	Sequence 6200	AA486628	dbEST
Sequence 6149	AA444386	dbEST	Sequence 6201	AA486837	dbEST
Sequence 6150	AA446103	dbEST	Sequence 6202	AA487192	dbEST
Sequence 6151	AA447349	dbEST	Sequence 6203	AA488014	dbEST
Sequence 6152	AA447563	dbES⊺	Sequence 6204	AA489323	dbEST
Sequence 6153	AA448526	dbEST	Sequence 6205	AA490047	dbEST
Sequence 6154	AA449228	dbEST	Sequence 6206	AA491272	dbEST
Sequence 6155	AA450120	dbEST	Sequence 6207	AA491302	dbEST
Sequence 6156	AA452177	dbEST	Sequence 6208	AA492026	dbEST
Sequence 6157	AA453134	dbEST	Sequence 6209	AA492038	dbEST
Sequence 6158	AA453310	dbEST	Sequence 6210	AA492042	dbEST
Sequence 6159	AA453333	dbEST	Sequence 6211	AA492219	dbEST
Sequence 6160	AA453478	dbEST	Sequence 6212	AA492280	dbEST
Sequence 6161	AA453719	dbEST	Sequence 6213	AA492390	dbEST
Sequence 6162	AA454962	dbEST	Sequence 6214	AA496932	dbEST
Sequence 6163	AA456246	dbEST	Sequence 6215	AA501968	dbEST
Sequence 6164	AA459208	dbEST	Sequence 6216	AA502897	dbEST
Sequence 6165	AA459871	dbEST	Sequence 6217	AA502955	dbEST
Sequence 6166	AA460440	dbEST	Sequence 6218	AA502979	dbEST
Sequence 6167	AA463469	dbEST	Sequence 6219	AA503718	dbEST
Sequence 6168	AA464160	dbEST	Sequence 6220	AA503930	dbEST
Sequence 6169	AA464338	dbEST	Sequence 6221	AA503943	dbEST
Sequence 6170	AA465494	dbEST	Sequence 6222	AA504173	dbEST
Sequence 6171	AA467748	dbEST	Sequence 6223	AA504246	dbEST
Sequence 6172	AA467761	dbEST	Sequence 6224	AA506607	dbEST
Sequence 6173	AA468151	dbEST	Sequence 6225	AA506923	dbEST
Sequence 6174	AA468167	dbEST	Sequence 6226	AA506967	dbEST
Sequence 6175	AA468504	dbEST	Sequence 6227	AA507084	dbEST dbEST
Sequence 6176	AA468760	dbEST	Sequence 6228	AA507511	
Sequence 6177	AA469129	dbEST	Sequence 6229	AA508552	dbEST
Sequence 6178	AA469151	dbEST	Sequence 6230	AA508579	dbEST

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Sequence 6231	AA508898	dbEST	Sequence 6283	AA574288	dbEST
Sequence 6232	AA513056	dbEST	Sequence 6284	AA574397	dbEST
Sequence 6233	AA513597	dbEST	Sequence 6285	AA574436	dbEST
Sequence 6234	AA514042	dbEST	Sequence 6286	AA576243	dbEST
Sequence 6235	AA515576	dbEST	Sequence 6287	AA576407	dbEST
Sequence 6236	AA515857	dbEST	Sequence 6288	AA578009	dbEST
Sequence 6237	AA516531	dbEST	Sequence 6289	AA578577	doEST
Sequence 6238	AA521140	dbEST	Sequence 6290	AA578646	dbEST
Sequence 6239	AA523252	dbEST	Sequence 6291	AA578773	dbEST
Sequence 6240	AA523671	dbEST	Sequence 6292	AA579039	dbEST
Sequence 6241	AA523902	dbEST	Sequence 6293	AA579904	dbEST
Sequence 6242	AA524651	dbEST	Sequence 6294	AA580097	dbEST
Sequence 6243	AA525032	dbEST	Sequence 6295	AA580829	dbEST
Sequence 6244	AA525419	dbEST	Sequence 6296	AA581716	dbEST
Sequence 6245	AA525497	dbEST	Sequence 6297	AA583350	dbEST
Sequence 6246	AA525958	dbEST	Sequence 6298	AA583353	dbEST
Sequence 6247	AA526244	dbEST	Sequence 6299	AA587020	dbEST
Sequence 6248	AA526283	dbEST	Sequence 6300	AA587243	dbEST
Sequence 6249	AA526834	dbEST	Sequence 6301	AA587459	dbEST
Sequence 6250	AA527667	dbEST	Sequence 6302	AA588685	dbEST
Sequence 6251	AA527805	dbEST	Sequence 6303	AA588754	dbEST
Sequence 6252	AA528204	dbEST	Sequence 6304	AA588882	dbEST
Sequence 6253	AA531198	dbEST	Sequence 6305	AA593159	dbEST
Sequence 6254	AA531509	dbEST	Sequence 6306	AA593668	dbEST
Sequence 6255	AA533115	dbEST	Sequence 6307	AA595100	dbEST
Sequence 6256	AA533386	dbEST	Sequence 6308	AA599416	dbEST
Sequence 6257	AA533494	dbEST	Sequence 6309	AA600257	dbEST
Sequence 6258	<b>AA533</b> 709	dbEST	Sequence 5310	AA603265	dbEST
Sequence 6259	<b>AA533</b> 785	dbEST	Sequence 6311	AA603834	dbEST
Sequence 6260	AA534246	dbEST	Sequence 6312	AA613908	dbEST
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Sequence 6263	AA535384	dbEST	Sequence 6315	AA621031	dbEST
Sequence 6264	<b>AA535</b> 695	dbEST	Sequence 6316	AA621073	dbEST
Sequence 6265	AA541677	dbEST	Sequence 6317	AA621820	dbEST
Sequence 6266	AA548308	dbEST	Sequence 6318	AA625415	dbEST
Sequence 6267	AA550927	dbEST	Sequence 6319	AA625485	dbEST
Sequence 6268	AA551894	dbEST	Sequence 6320	AA625833	dbEST
Sequence 6269	AA552321	dbEST	Sequence 6321	AA626161	dbEST
Sequence 6270	AA552842	dbEST	Sequence 6322	AA626274	dbEST
Sequence 6271	AA558611	dbEST	Sequence 6323	AA626439	dbEST
Sequence 6272	AA558636	dbEST	Sequence 6324	AA628530	dbEST
Sequence 6273	AA558976	dbEST	Sequence 6325	AA629325	dbEST
Sequence 6274	AA564277	dbEST	Sequence 6326	AA629581	dbEST
Sequence 6275	<b>AA5</b> 64484	dbEST	Sequence 6327	AA630298	dbEST
Sequence 6276	AA565996	dbEST	Sequence 6328	AA631446	dbEST
Sequence 6277	AA569344	dbEST	Sequence 6329	AA631868	dbEST
Sequence 6278	AA569434	dbEST	Sequence 6330	AA632427	dbEST
Sequence 6279	AA569482	dbEST	Sequence 6331	AA633399	dbEST
Sequence 6280	AA573368	dbEST	Sequence 6332	AA633846	dbEST
Sequence 6281	AA573646	dbEST	Sequence 6333	AA635288	cbEST
Sequence 6282	AA573742	dbEST	Sequence 6334	AA640133	cbEST

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Sequence 6335	AA640574	dbEST	Sequence 6387	AA758592	dbEST
Sequence 6336	AA640889	dbEST	Sequence 6388	AA761739	dbEST
Sequence 6337	AA641585	dbEST	Sequence 6389	AA768762	dbEST
Sequence 6338	AA641623	dbEST	Sequence 6390	AA774265	dbEST
Sequence 6339	AA642495	dbEST	Sequence 6391	AA775625	dbEST
Sequence 6340	AA648194	dbEST	Sequence 6392	AA776709	dbEST
Sequence 6341	AA652478	dbEST	Sequence 6393	AA779427	dbEST
Sequence 6342	AA654527	dbEST	Sequence 6394	AA805361	dbEST
Sequence 6343	AA657830	dbEST	Sequence 6395	AA806555	dbEST
Sequence 6344	AA659142	dbEST	Sequence 6396	AA813373	dbEST
Sequence 6345	AA659391	dbEST	Sequence 6397	AA814822	dbEST
Sequence 6346	AA662111	dbEST	Sequence 6398	AA824416	dbEST
Sequence 6347	AA662369	dbEST	Sequence 6399	AA825612	dbEST
Sequence 6348	AA662822	dbEST	Sequence 6400	AA827079	dbEST
Sequence 6349	AA663453	dbEST	Sequence 6401	AA827968	dbEST
Sequence 6350	AA664480	dbEST	Sequence 6402	AA828243	dbEST
Sequence 6351	AA669438	dbEST	Sequence 6403	AA830856	dbEST
Sequence 6352	AA670205	dbEST	Sequence 6404	AA831148	dbEST
Sequence 6353	AA677550	dbEST	Sequence 6405	AA835638	dbEST
Sequence 6354	AA687179	dbEST	Sequence 6406	AA838377	dbEST
Sequence 6355	AA687216	dbEST	Sequence 6407	AA838431	dbEST
Sequence 6356	AA687370	dbEST	Sequence 6408	AA838482	dbEST
Sequence 6357	AA687538	dbEST	Sequence 6409	AA843133	dbEST
Sequence 6358	AA687674	dbEST	Sequence 6410	AA845390	dbEST
Sequence 6359	AA687718	dbEST	Sequence 6411	AA846335	dbEST
Sequence 6360	AA693742	dbEST	Sequence 6412	AA852998	dbEST
Sequence 6361	AA694055	dbEST	Sequence 6413	AA857872	dbEST
Sequence 6362	AA694120	dbEST	Sequence 6414	AA861986	dbEST
Sequence 6363	AA700509	dbEST	Sequence 6415	AA862309	dbEST
Sequence 6364	AA701607	dbEST	Sequence 6416	AA864840	dbEST
Sequence 6365	AA702084	dbEST	Sequence 6417	AA872710	dbEST
Sequence 6366	AA702131	dbEST	Sequence 6418	AA873534	dbEST
Sequence 6367	AA702891	dbEST	Sequence 6419	AA877864	dbEST
Sequence 6368	AA703838	dbEST	Sequence 6420	AA883508	dbEST
Sequence 6369	AA703925	dbEST	Sequence 6421	AA885780	dbEST
Sequence 6370	AA704323	dbEST	Sequence 6422	AA885835	dbEST
Sequence 6371	AA705957	dbEST	Sequence 6423	AA887235	dbEST
Sequence 6372	AA705986	dbEST	Sequence 6424	AA890032	dbEST
Sequence 6373	AA706314	dbEST	Sequence 6425	AA894781	dbEST
Sequence 6374	AA707377	dbEST	Sequence 6426	AA894929	dbEST
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Sequence 6379	AA716097	dbEST	Sequence 6431	AA905065	dbEST
Sequence 6380	AA719530	dbEST	Sequence 6432	AA908930	dbEST
Sequence 6381	AA725298	dbEST	Sequence 6433	AA909977	dbEST
Sequence 6382	AA725836	dbEST	Sequence 6434	AA911903	dbEST
Sequence 6383	AA732352	dbEST	Sequence 6435	AA912713	dbEST
Sequence 6384	AA745678	dbEST	Sequence 6436	AA917937	dbEST
Sequence 6385	AA747515	dbEST	Sequence 6437	AA923547	dbEST
Sequence 6386	AA748033	dbEST	Sequence 6438	AA926656	dbEST

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Sequence 6439	AA928427	dbEST	Sequence 6491	AI085393	dbEST
Sequence 6440	AA932331	dbEST	Sequence 6492	AI087412	dbEST
Sequence 6441	AA934851	dbEST	Sequence 6493	AI089394	dbEST
Sequence 6442	AA935485	dbEST	Sequence 6494	AI089526	dbEST
Sequence 6443	AA935560	dbEST	Sequence 6495	AI089948	dbEST
Sequence 6444	AA936089	dbEST	Sequence 6496	AI090786	dbEST
Sequence 6445	AA937513	dbEST	Sequence 6497	AI091736	dbEST
Sequence 6446	AA937773	dbEST	Sequence 6498	AI095875	dbEST
Sequence 6447	AA938181	dbEST	Sequence 6499	Al096493	dbEST
Sequence 6448	AA938717	dbEST	Sequence 6500	Al123916	dbEST
Sequence 6449	AA939023	dbEST	Sequence 6501	AI124532	dbEST
Sequence 6450	AA948276	dbEST	Sequence 6502	Al125642	dbEST
Sequence 6451	AA970266	dbEST	Sequence 6503	AI127167	dbEST
Sequence 6452	AA972530	dbEST	Sequence 6504	AI127176	dbEST
Sequence 6453	AA972602	dbEST	Sequence 6505	Al 127424	dbEST
Sequence 6454	AA974137	dbEST	Sequence 6506	AJ127455	dbEST
Sequence 6455	AA975275	dbEST	Sequence 6507	Al127556	dbEST
Sequence 6456	AA977226	dbEST	Sequence 6508	AI128918	dbEST
Sequence 6457	AA977833	dbEST	Sequence 6509	Al129342	dbEST
Sequence 6458	AA978353	dbEST	Sequence 6510	Al129403	dbEST
Sequence 6459	AA983560	dbEST	Sequence 6511	Al133096	dbEST
Sequence 6460	AA985190	dbEST	Sequence 6512	Al133152	dbEST
Sequence 6461	AA994534	dbEST	Sequence 6513	AI133330	dbEST
Sequence 6462	AA995483	dbEST	Sequence 6514	AI133381	dbEST
Sequence 6463	AF017688	dbEST	Sequence 6515	Al133402	dbEST
Sequence 6464	AF034176	dbEST	Sequence 6516	Al139036	dbEST
Sequence 6465	AI015053	dbEST	Sequence 6517	AI139157	dbEST
Sequence 6466	AI016700	dbEST	Sequence 6518	Al140251	dbEST
Sequence 6467	AI018625	dbEST	Sequence 6519	Al140421	dbEST
Sequence 6468	Al022134	dbEST	Sequence 6520	Al141483	dbEST
Sequence 6469	AI022855	dbEST	Sequence 6521	Al142083	dbEST
Sequence 6470	AI023931	dbEST	Sequence 6522	Al148437	dbEST
Sequence 6471	AI027887	dbEST	Sequence 6523	Al149743	dbEST
Sequence 6472	Al028122	dbEST	Sequence 6524	Al151482	dbEST
Sequence 6473	AI028708	dbEST	Sequence 6525	AI160324	dbEST
Sequence 6474	AI032401	dbEST	Sequence 6526	AI161333	dbEST
Sequence 6475	AI034431	dbEST	Sequence 6527	Al174828	dbEST
Sequence 6476	AI041847	dbEST	Sequence 6528	Al186545	dbEST
Sequence 6477	AI042059	dbEST	Sequence 6529	Al188641	dbEST
Sequence 6478	AI052112	dbEST	Sequence 6530	Al190480	dbEST
Sequence 6479	AI055870	dbEST	Sequence 6531	Al192932	dbEST
Sequence 6480	AI056489	dbEST	Sequence 6532	AI197946	dbEST
Sequence 6481	Al057142	dbEST	Sequence 6533	AI198625	dbEST
Sequence 6482	AJ074397	dbEST	Sequence 6534	Al199544	dbEST
Sequence 6483	AI074613	dbEST	Sequence 6535	AI200798	dbEST
Sequence 6484	AI078381	dbEST	Sequence 6536	AI202026	dbEST
Sequence 6485	AI080170	dbEST	Sequence 6537	AI203141	dbEST
Sequence 6486	A1080267	dbEST	Sequence 6538	Al204308	dbEST
Sequence 6487	AI080485	dbEST	Sequence 6539	Al205228	dbEST
Sequence 6488	AI081291	dbEST	Sequence 6540	Al206063	dbEST
Sequence 6489	AI084101	dbEST	Sequence 6541	AI207781	dbEST
Sequence 6490	AI084916	dbEST	Sequence 6542	AI216969	dbEST

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Sequence 6543	Al217009	dbEST	Sequence 6595	Al347473	dbEST
Sequence 6544	AJ217172	dbEST	Sequence 6596	Al349099	dbEST
Sequence 6545	Al217964	dbEST	Sequence 6597	AI34 <b>9937</b>	dbEST
Sequence 6546	Al219278	dbEST	Sequence 6598	Al357472	dbEST
Sequence 6547	Al220051	dbEST	Sequence 6599	AI360165	dbEST
Sequence 6548	Al239506	dbEST	Sequence 6600	A1366549	dbEST
Sequence 6549	Al241117	dbEST	Sequence 6601	Al368743	dbEST
Sequence 6550	Al241659	dbEST	Sequence 6602	Al369849	dbEST
Sequence 6551	Al241744	dbEST	Sequence 6603	Al377012	dbEST
Sequence 6552	AI246383	dbE <b>ST</b>	Sequence 6604	Al378613	dbEST
Sequence 6553	Al249764	dbEST	Sequence 6605	Al380537	dbEST
Sequence 6554	A1250055	dbE <b>S</b> T	Sequence 6606	Al380842	dbEST
Sequence 6555	Al252084	dbEST	Sequence 6607	A1401771	dbEST
Sequence 6556	Al252283	dbEST	Sequence 6608	Al417016	dbEST
Sequence 6557	Al253099	dbEST	Sequence 6609	Al421249	dbEST
Sequence 6558	AI253330	dbEST	Sequence 6610	AI431350	dbEST
Sequence 6559	Al253335	dbEST	Sequence 6611	Al433818	dbEST
Sequence 6560	Al253363	dbEST	Sequence 6612	Al434250	dbEST
Sequence 6561	Al261763	dbEST	Sequence 6613	AI435349	dbEST
Sequence 6562	Al267185	dbEST	Sequence 6614	A1439987	dbEST
Sequence 6563	Al267282	dbEST	Sequence 6615	Al446249	dbEST
Sequence 6564	Al267502	dbEST	Sequence 6616	AI452707	dbEST
Sequence 6565	Al269158	dbEST	Sequence 6617	Al453402	dbEST
Sequence 6566	Al273636	dbEST	Sequence 6618	Al474125	dbEST
Sequence 6567	Al276399	dbEST	Sequence 6619	Al475810	dbEST
Secuence 6568	Al276576	dbEST	Sequence 6620	A1479948	dbEST
Sequence 6569	Al276662	dbEST	Sequence 6621	Al491775	dbEST
Sequence 6570	Al278014	dbEST	Sequence 6622	Al499331	dbEST
Sequence 6571	Al281196	dbEST	Sequence 6623	A1525654	dbEST
Sequence 6572	Al281926	dbEST	Sequence 6624	AI525843	dbEST
Sequence 6573	Al282318	dbEST	Sequence 6625	AI536833	dbEST
Sequence 6574	Al298059	dbEST	Sequence 6626	Al547041	dbEST
Sequence 6575	Al300532	dbEST	Sequence 6627	A1547068	dbEST
Sequence 6576	Al300553	dbEST	Sequence 6628	A1547090	dbEST
Sequence 6577	Al302799	dbEST	Sequence 6629	AI547125	dbEST
Sequence 6578	Al304935	dbEST	Sequence 6630	A1547285	dbEST
Sequence 6579	Al310138	dbEST	Sequence 6631	AI557225	dbEST
Sequence 6580	Al312325	dbEST	Sequence 6632	AI557231	dbEST
Sequence 6581	Al318091	dbEST	Sequence 6633	AI557246	dbEST
Sequence 6582	Al335028	dbEST	Sequence 6634	A1557269	dbEST
Sequence 6583	Al335178	dbEST	Sequence 6635	AI557458	dbEST
Sequence 6584	Al336501	dbEST	Sequence 6636	AI557495	dbEST
Sequence 6585	Al336800	dbEST	Sequence 6637	AI563921	dbEST
Sequence 6586	Al336949	dbEST	Sequence 6638	AI566084	dbEST
Sequence 6587	Al338603	dbEST	Sequence 6639	AI570349	dbEST
Sequence 6588	Al339354	dbEST	Sequence 6640	AI612873	dbEST
Sequence 6589	Al340981	dbEST	Sequence 6641	AI627356	dbEST
Sequence 6590	Al341972	dbEST	Sequence 6642	AI630118	dbEST
Sequence 6591	Al342546	dbEST	Sequence 6643	AI630685	dbEST
Sequence 6592	Al342937	dbEST	Sequence 6644	AI630985	dbEST
Sequence 6593	Al343692	dbEST	Sequence 6645	AI631745	dbEST
Sequence 6594	Al346657	dbEST	Sequence 6646	Al632244	dbEST

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Sequence		AJ632534	dbEST	Sequence 6699	Al929339	dbEST
Sequence		AI636701	db <b>EST</b>	Sequence 6700	Al934595	dbEST
Sequence		AI653367	dbEST	Sequence 6701	Al951049	dbEST
Sequence	6650	Al654122	dbEST	Sequence 6702	AI952286	dbEST
Sequence	6651	AI6598 <b>98</b>	dbEST	Sequence 6703	Al961413	dbEST
Sequence	6652	Al660243	dbEST	Sequence 6704	Al968069	dbEST
Sequence	6653	AI669052	dbEST	Sequence 6705	A1969804	dbEST
Sequence	6654	Al674586	dbEST	Sequence 6706	A1978832	dbEST
Sequence	6655	Al680416	d <b>bEST</b>	Sequence 6707	AL035756	dbEST
Sequence	6656	A1680506	d <b>bEST</b>	Sequence 6708	AL036801	dbEST
Sequence	6657	Al684170	dbEST	Sequence 6709	AL036890	dbEST
Sequence	6658	Al691072	dbEST	Sequence 6710	AL037798	dbEST
Sequence	6659	Al696513	dbEST	Sequence 6711	AL038242	dbEST
Sequence	6660	AI700228	d <b>bEST</b>	Sequence 6712	AL038395	dbEST
Sequence	6661	Al719425	dbEST	Sequence 6713	AL039175	dbEST
Sequence	6662	Al719701	d <b>bEST</b>	Sequence 6714	AL039507	dbEST
Sequence	6663	AI720354	dbEST	Sequence 6715	AL040632	dbEST
Sequence	6664	AI732097	dbEST	Sequence 6716	AL040676	dbEST
Sequence	6665	AI734266	dbEST	Sequence 6717	AL044040	dbEST
Sequence	6666	Al734854	dbEST	Sequence 6718	AL045977	dbEST
Sequence	6667	A1735683	dbEST	Sequence 6719	AL046005	dbEST
Sequence	6668	AI740869	dbEST	Sequence 6720	AL046079	dbEST
Sequence	6669	AI741192	dbEST	Sequence 6721	AL048748	dbEST
Sequence	6670	AI749547	dbEST	Sequence 6722	AL049145	dbEST
Sequenc <b>e</b>	6671	AI751930	dbEST	Sequence 6723	AL118746	dbEST
Sequence	6672	AI752546	dbEST	Sequence 6724	AL119449	dbEST
Sequence	6673	Al752913	dbEST	Sequence 6725	AL119462	dbEST
Sequence	6674	Al753229	dbEST	Sequence 6726	AL120558	dbEST
Sequence		AI753662	dbEST	Sequence 6727	AL120741	dbEST
Sequence		AI753951	dbEST	Sequence 6728	AL120789	dbEST
Sequence		AI754071	dbEST	Sequence 6729	AL121220	dbEST
Sequence		AI755085	dbEST	Sequence 6730	AL121488	dbEST
Sequence		Al796120	dbEST	Sequence 6731	AW003821	dbEST
Sequence		Al810698	dbEST	Sequence 6732	AW003952	dbEST
Sequence		AI813937	dbEST	Sequence 6733	AW005947	dbEST
Sequence		AI815677	dbEST	Sequence 6734	AW015229	dbEST
Sequence		AI815703	dbEST	Sequence 6735	AW015828	dbEST
Sequence		Al815972	dbEST	Sequence 6736	AW021832	dbEST
Sequence		AI815987	dbEST	Sequence 6737	AW026131	dbEST
Sequence		AI816235	dbEST	Sequence 6738	AW026502	dbEST
Sequence		AI816313	dbEST	Sequence 6739	AW055209	dbEST
Sequence		AI816938	dbEST	Sequence 6740	AW068372	dbEST
Sequence		AI818255	dbEST	Sequence 6741	AW082236	dbEST
Sequence		A1825504	dbEST	Sequence 6742	AW087623	dbEST
Sequence		Al825799	dbEST	Sequence 6743	AW090304	dbEST
Sequence		AI859156	dbEST	Sequence 6744	AW102895	dbEST
Sequence		A1878846	dbEST	Sequence 6745	AW130969	dbEST
Sequence		AI879036	dbEST	Sequence 6746	AW135062	dbEST
Sequence		AI884963	dbEST	Sequence 6747	AW135187	dbEST
Sequence		AI889347	dbEST	Sequence 6748	AW157303	dbEST
Sequence		AI914133	dbEST	Sequence 6749	AW160399	dbEST
Sequence	0098	Al917110	dbEST	Sequence 6750	AW160696	dbEST

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C	A14404000	AFECT	Camuanaa 6903	N79712	dbEST
Sequence 6751	AW161209	dbEST	Sequence 6803 Sequence 6804	N94351	dbEST
Sequence 6752	AW161365	dbEST	•	N98328	dbEST
Sequence 6753	AW161467	dbEST	Sequence 6805		
Sequence 6754	AW162502	dbEST	Sequence 6806	N99088	dbEST
Sequence 6755	AW176042	dbEST	Sequence 6807	R12356	dbEST
Sequence 6756	AW178342	dbEST	Sequence 6808	R16295	dhEST
Sequence 6757	AW179006	dbEST	Sequence 6809	R16408	dbEST
Sequence 6758	C03130	dbEST	Sequence 6810	R19453	dbEST
Sequence 6759	C18148	dbEST	Sequence 6811	R19869	dbEST
Sequence 6760	C18654	dbEST	Sequence 6812	R20063	dbEST
Sequence 6761	D12135	dbEST	Sequence 6813	R24400	dbEST
Sequence 6762	D30921	dbEST	Sequence 6814	R25464	dbEST
Sequence 6763	D58694	dbEST	Sequence 6815	R33351	dbEST
Sequence 6764	D61285	dbEST	Sequence 6816	R45686	dbEST
Sequence 6765	D82149	dbEST	Sequence 6817	R53758	dbEST
Sequence 6766	D82176	dbEST	Sequence 6818	R56382	dbEST
Sequence 6767	F01265	dbEST	Sequence 6819	R61196	dbEST
Sequence 6768	F01327	dbEST	Sequence 6820	R661 <b>62</b>	dbEST
Sequence 6769	F05508	dbEST	Sequence 6821	R70193	dbEST
Sequence 6770	H02725	dbEST	Sequence 6822	R70570	dbEST
Sequence 6771	H04072	dbEST	Sequence 6823	R72768	dbEST
Sequence 6772	H10429	dbEST	Sequence 6824	R91218	dbEST
Sequence 6773	H10482	dbEST	Sequence 6825	R94399	dbEST
Sequence 6774	H12109	dbEST	Sequence 6826	R99755	dbEST
Sequence 6775	H12519	dbEST	Sequence 6827	T08329	dbEST
Sequence 6776	H13339	dbEST	Sequence 6828	T34459	dbEST
Sequence 6777	H17773	dbEST	Sequence 6829	T35276	dbEST
Sequence 6778	H19393	dbEST	Sequence 6830	T60601	dbEST
Sequence 6779	H38564	dbEST	Sequence 6831	T67978	dbEST
Sequence 6780	H47325	dbEST	Sequence 6832	T68894	dbEST
Sequence 6781	H48697	dbEST	Sequence 6833	T71387	dbEST
Sequence 6782	H51749	dbEST	Sequence 6834	T73262	dbEST
Sequence 6783	H54240	dbEST	Sequence 6835	T81537	dbEST
Sequence 6784	H59731	dbEST	Sequence 6836	W15207	dbEST
Sequence 6785	H66398	dbEST	Sequence 6837	W15554	dbEST
Sequence 6786		dbEST	Sequence 6838	W25142	dbEST
Sequence 6787	H84670	dbEST	Sequence 6839	W26372	dbEST
Sequence 6788		dbEST	Sequence 6840	W30832	dbEST
Sequence 6789		dbEST	Sequence 6841	W38726	dbEST
Sequence 6790		dbEST	Sequence 6842	W38 <b>956</b>	dbEST
Sequence 6791	N23771	dbEST	Sequence 6843	W42913	dbEST
Sequence 6792	N39674	dbEST	Sequence 6844	W73290	dbEST
		dbEST	Sequence 6845	W76133	dbEST
Sequence 6793	N39836 N40375	dbEST	Sequence 6846	W80496	dbEST
Sequence 6794			Sequence 6847	W81084	dbEST
Sequence 6795		dbEST	Sequence 6848	N40101	NUCPATENT
Sequence 6796		dbEST	Sequence 6849	N91258	NUCPATENT
Sequence 6797		dbEST		Q24595	NUCPATENT
Sequence 6798		dbEST	Sequence 6850	Q63815	NUCPATENT
Sequence 6799		dbEST	Sequence 6851	Q97381	NUCPATENT
Sequence 6800		dbEST	Sequence 6852	T74203	NUCPATENT
Sequence 6801	N63387	dbEST	Sequence 6853	V07906	NUCPATENT
Sequence 6802	N75468	dbEST	Sequence 6854	V0/900	HOULAILIN

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Sequence 6855	V30916	NUCPATENT	Sequence 6905	AF035839	GENBANK
Sequence 6856	V34207	NUCPATENT	Sequence 6906	AF044957	GENBANK
Sequence 6857	V34226	NUCPATENT	Sequence 6907	AF048977	<b>GENBANK</b>
Sequence 6858	V34305	NUCPATENT	Sequence 6908	AF052138	GENBANK
Sequence 6859	V35555	NUCPATENT	Sequence 6909	AF054183	GENBANK
Sequence 6860	V43604	NUCPATENT	Sequence 6910	AF054987	GENBANK
Sequence 6861	V58584	NUCPATENT	Sequence 6911	AF055473	GENBANK
Sequence 6862	V58756	NUCPATENT	Sequence 6912	AF055474	GENBANK
Sequence 6863	V58761	NUCPATENT	Sequence 6913	AF067396	GENBANK
Sequence 6864	V62427	NUCPATENT	Sequence 6914	AF068117	GENBANK
Sequence 6865	V69855	NUCPATENT	Sequence 6915	AF068754	GENBANK
Sequence 6866	V70354	NUCPATENT	Sequence 6916	AF069301	GENBANK
Sequence 6867	V72125	NUCPATENT	Sequence 6917	AF070655	GENBANK
Sequence 6868	V72211	NUCPATENT	Sequence 6918	AF071172	GENBANK
Sequence 6869	V84448	NUCPATENT	Sequence 6919	AF071748	GENBANK
Sequence 6870	X27262	NUCPATENT	Sequence 6920	AF073298	GENBANK
Sequence 6871	X27278	NUCPATENT	Sequence 6921	AF077045	GENBANK
Sequence 6872	X30167	NUCPATENT			
Sequence 6873	X36344	NUCPATENT	Sequence 6922	AF077202 AF083190	GENBANK
Sequence 6874	X36833	NUCPATENT	Sequence 6923		GENBANK
Sequence 6875	X37486	NUCPATENT	Sequence 6924	AF085844	GENBANK
Sequence 6876	X39662	NUCPATENT	Sequence 6925	AF086251	GENBANK
Sequence 6877	X40462	NUCPATENT	Sequence 6926	AF088035	GENBANK
Sequence 6878	X51602	NUCPATENT	Sequence 6927	AF091073	GENBANK
Sequence 6879	X97772	NUCPATENT	Sequence 6928	AF100759	GENBANK
Sequence 6880	X97841	NUCPATENT	Sequence 6929	AF101044	GENBANK
Sequence 6881	X97888	NUCPATENT	Sequence 6930	AF140598	GENBANK
Sequence 6882	X98901	NUCPATENT	Sequence 6931	AF147330	GENBANK
Sequence 6883	X99098	NUCPATENT	Sequence 6932	AF151823	GENBANK
Sequence 6884	Z00842	NUCPATENT	Sequence 6933	AF151830	GENBANK
Sequence 6885	Z06250	NUCPATENT	Sequence 6934	AF153603	GENBANK
Sequence 6886	Z10658	NUCPATENT	Sequence 6935	AF153612	GENBANK
Sequence 6887	Z15890	NUCPATENT	Sequence 6936	AF170708	GENBANK
Sequence 6888	Z24862	NUCPATENT	Sequence 6937	AF176574	GENBANK
Sequence 6889	AC29871	PREPATNUC	Sequence 6938	AJ007398	GENBANK
Sequence 6890	AC31081	PREPATNUC	Sequence 6939	AL049981	GENBANK
Sequence 6891	AC33416	PREPATNUC	Sequence 6940	AL050089	GENBANK
Sequence 6892	AC33722	PREPATNUC	Sequence 6941	AL050291	GENBANK
Sequence 6893	AC33900	PREPATNUC	Sequence 6942	AL080089	GENBANK
Sequence 6894	AC34235	PREPATNUC	Sequence 6943	AL110185	GENBANK
Sequence 6895	AC34354	PREPATNUC	Sequence 6944	AL110191	GENBANK
Sequence 6896	AC34866	PREPATNUC	Sequence 6945	AL122063	GENBANK
Sequence 6897	AC34877	PREPATNUC	Sequence 6946	D14697	GENBANK
			Sequence 6947	D16469	GENBANK
Sequence	Accession	# Database	Sequence 6948	D45248	GENBANK
Sequence 6898	AB000468	GENBANK	Sequence 6949	D49396	GENBANK
Sequence 6899	AB004788	GENBANK	Sequence 6950	D50371	GENBANK
Sequence 6900	AB007893	GENBANK	Sequence 6951	D50525	GENBANK
Sequence 6901	AB009285	GENBANK	Sequence 6952	D50916	GENBANK
Sequence 6902	AB015644	GENBANK	Sequence 6953	D876 <b>66</b>	GENBANK
Sequence 6903	AB016087	GENBANK	Sequence 6954	D88153	GENBANK
Sequence 6904	AF020351	GENBANK	Sequence 6955	E01915	GENBANK
		CE. 10/ 11/11			

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Sequence 6956	J02642	GENBANK	Sequence 7007	AA042813	dbEST
Sequence 6957	J03779	GENBANK	Sequence 7008	AA044697	dbEST
Sequence 6958	J04208	GENBANK	Sequence 7009	AA045753	dbEST
Sequence 6959	L03558	GENBANK	Sequence 7010	AA045879	dbEST
Sequence 6960	L050 <b>93</b>	GENBANK	Sequence 7011	AA057541	dbEST
Sequence 6961	L14778	GENBANK	Sequence 7012	AA062804	dbEST
Sequence 6962	L20422	GENBANK	Sequence 7013	AA062928	dbEST
Sequence 6963	M11147	GENBANK	Sequence 7014	AA064627	dbEST
Sequence 6964	M11560	GENBANK	Sequence 7015	AA070911	dbEST
Sequence 6965	M17885	GENBANK	Sequence 7016	AA076077	dbEST
Sequence 6966	M37583	GENBANK	Sequence 7017	AA076593	dbEST
Sequence 6967	M38188	GENBANK	Sequence 7018	AA081815	dbEST
Sequence 6968	M64241	GENBANK	Sequence 7019	AA082502	dbEST
Sequence 6969	U12979	GENBANK	Sequence 7020	AA083481	dbEST
Sequence 6970	U17104	GENBANK	Sequence 7021	AA083794	dbEST
Sequence 6971	U17899	GENBANK	Sequence 7022	AA083868	dbEST
Sequence 6972	U19143	GENBANK	Sequence 7023	AA085603	dbEST
Sequence 6973	U19144	GENBANK	Sequence 7024	AA085824	dbEST
Sequence 6974	U22431	GENBANK	Sequence 7025	AA094583	dbEST
Sequence 6975	U73824	GENBANK	Sequence 7026	AA095753	dbEST
Sequence 6976	U75330	GENBANK	Sequence 7027	AA098801	dbEST
Sequence 6977	X07767	GENBANK	Sequence 7028	AA099023	dbEST
Sequence 6978	X62048	GENBANK	Sequence 7029	AA100987	dbEST
Sequence 6979	X63423	GENBANK	Sequence 7030	AA121390	dbEST
Sequence 6980	X75535	GENBANK	Sequence 7031	AA127493	dbEST
Sequence 6981	X89673	GENBANK	Sequence 7032	AA128952	dbEST
Sequence 6982	X99585	GENBANK	Sequence 7033	AA129726	dbEST
Sequence 6983	Z11692	GENBANK	Sequence 7034	AA130873	dbEST
Sequence 6984	Z19002	GENBANK	Sequence 7035	AA132219	dbEST
Sequence 6985	Z47087	GENBANK	Sequence 7036	AA132574	dbEST
Sequence 6986	Z48054	GENBANK	Sequence 7037	AA132690	dbEST
Sequence 6987	AA001699	dbEST	Sequence 7038	AA133228	dbEST
Sequence 6988	AA004206	dbEST	Sequence 7039	AA133266	dbEST
Sequence 698	AA005403	dbEST	Sequence 7040	AA135049	dbEST
Sequence 6990	AA009849	dbEST	Sequence 7041	AA136150	dbEST
Sequence 6991	AA010129	dbEST	Sequence 7042	AA136296	dbEST
Sequence 6992	AA016205	dbEST	Sequence 7043	AA147986	dbEST
Sequence 6993	AA018050	dbEST	Sequence 7044	AA149080	dbEST
Sequence 6994	AA025694	dbEST	Sequence 7045	AA150156	dbEST
Sequence 6995	AA027090	dbEST	Sequence 7046	AA150267	dbEST
Sequence 6996	AA029566	dbEST	Sequence 7047	AA150369	dbEST
Sequence 6997	AA031624	dbEST	Sequence 7048	AA150839	dbEST
Sequence 6998	AA031795	dbEST	Sequence 7049	AA150875	dbEST
Sequence 6999	AA033537	dbEST	Sequence 7050	AA152408	dbEST
Sequence 7000	AA039274	dbEST	Sequence 7051	AA156743	dbEST
Sequence 7001	AA039506	dbEST	Sequence 7052	AA157816	dbEST
Sequence 7002	AA039734	dbEST	Sequence 7053	AA158098	dbEST
Sequence 7003	AA039750	dbEST	Sequence 7054	AA159044	dbEST
Sequence 7004	AA039807	dbEST	Sequence 7055	AA160380	dbEST
Sequence 7005	AA040499	dbEST	Sequence 7056	AA164405	dbEST
Sequence 7006	AA040625	dbEST	Sequence 7057	AA165117	dbEST
			•		

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Sequence 7058	AA166853	dbEST	Sequence 7109	AA328061	dbEST
Sequence 7059	AA167011	dbEST	Sequence 7110	AA329553	dbEST
Sequence 7060	AA169493	dbEST	Sequence 7111	AA351201	dbEST
Sequence 7061	AA171618	dbEST	Sequence 7112	AA354420	dbEST
Sequence 7062	AA171706	dbEST	Sequence 7113	AA380769	dbEST
Sequence 7063	AA177062	dbEST	Sequence 7/114	AA393336	dbEST
Sequence 7064	AA180137	dbEST	Sequence 7115	AA398256	dbEST
Sequence 7065	AA180817	dbEST	Sequence 7116	AA399238	dbEST
Sequence 7066	AA191045	dbEST	Sequence 7117	AA400658	· dbEST
Sequence 7067	AA194429	dbEST	Sequence 7118	AA401237	dbEST
Sequence 7068	AA194535	dbEST	Sequence 7119	AA402330	dbEST
Sequence 7069	AA195470	dbEST	Sequence 7120	AA404535	dbEST
Sequence 7070	AA203462	dbEST	Sequence 7121	AA405929	dbEST
Sequence 7071	AA204856	dbEST	Sequence 7122	AA406122	dbEST
Sequence 7072	AA205858	dbEST	Sequence 7123	AA406404	dbEST
Sequence 7073	AA216133	dbEST	Sequence 7124	AA406565	dbEST
Sequence 7074	AA216211	dbEST	Sequence 7125	AA410640	dbEST
Sequence 7075	AA219582	dbEST	Sequence 7126	AA412229	dbEST
Sequence 7076	AA226340	dbEST	Sequence 7127	AA420563	dbEST
Sequence 7077	AA227105	dbEST	Sequence 7128	AA424175	dbEST
Sequence 7078	AA227695	dbEST	Sequence 7129	AA425389	dbEST
Sequence 7079	AA229607	dbEST	Sequence 7130	AA429036	dbEST
Sequence 7080	AA229611	dbEST	Sequence 7131	AA429457	dbEST
Sequence 7081	AA232686	dbEST	Sequence 7132	AA431401	dbEST
Sequence 7082	AA233053	dbEST	Sequence 7133	AA435877	dbEST
Sequence 7083	AA253254	dbEST	Sequence 7134	AA443357	dbEST
Sequence 7084	AA255502	dbEST	Sequence 7135	AA447559	dbEST
Sequence 7085	AA255529	dbEST	Sequence 7136	AA448542	dbEST
Sequence 7086	AA256507	dbEST	Sequence 7137	AA449663	dbEST
Sequence 7087	AA280381	dbEST	Sequence 7138	AA449891	dbEST
Sequence 7088	AA282837	dbEST	Sequence 7139	AA452722	dbEST
Sequence 7089	AA284506	dbEST	Sequence 7140	AA459869	dbEST
Sequence 7090	AA285089	dbEST	Sequence 7141	AA460777	dbEST
Sequence 7091	AA285290	dbEST	Sequence 7142	AA469432	dbEST
Sequence 7092	AA287721	dbEST	Sequence 7143	AA470662	dbEST
Sequence 7093	AA292273	dbEST	Sequence 7144	AA470829	dbEST
Sequence 7094	AA296152	dbEST	Sequence 7145	AA477597	dbEST
Sequence 7095	AA296237	dbEST	Sequence 7146	AA479363	dbEST
Sequence 7096	AA301795	dbEST	Sequence 7147	AA480202	dbEST
Sequence 7097	AA306372	dbEST	Sequence 7148	AA480226	dbEST
Sequence 7098	AA306982	dbEST	Sequence 7149	AA480863	dbEST
Sequence 7099	AA307814	dbEST	Sequence 7150	AA483013	dbEST
Sequence 7100	AA307908	dbEST	Sequence 7151	AA483289	dbEST
Sequence 7101	AA313647	dbEST	Sequence 7152	AA483857	dbEST
Sequence 7102	AA313693	dbEST	Sequence 7153	AA485065	dbEST
Sequence 7103	AA314391	dbEST	Sequence 7154	AA487463	dbEST
Sequence 7104	AA316788	dbEST	Sequence 7155	AA489074	dbEST
Sequence 7105	AA316962	dbEST	Sequence 7156	AA492038	dbEST
Sequence 7106	AA318100	dbEST	Sequence 7157	AA501447	dbEST
Sequence 7107	AA318712	dbEST	Sequence 7158	AA502178	dbEST
Sequence 7108	AA325594	dbEST	Sequence 7159	AA502730	dbEST

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Sequence 7160	AA504514	dbEST	Sequence 7211	AA613314	dbEST
Sequence 7161	AA504622	dbEST	Sequence 7212	AA617696	dbEST
Sequence 7162	AA506953	dbEST	Sequence 7213	AA618183	dbEST
Sequence 7163	AA508507	dbEST	Sequence 7214	AA618435	dbEST
Sequence 7164	AA512975	dbEST	Sequence 7215	AA621730	dbEST
Sequence 7165	AA514324	dbEST	Sequence 7216	AA625270	dbEST
Sequence 7166	AA514474	dbEST	Sequence 7217	AA629269	dbEST
Sequence 7167	AA521013	dbEST	Sequence 7218	AA631009	dbEST
Sequence 7168	AA522638	dbEST	Sequence 7219	AA631608	dbEST
Sequence 7169	AA522746	dbEST	Sequence 7220	AA632154	dbEST
Sequence 7170	AA522849	dbEST	Sequence 7221	AA635393	dbEST
Sequence 7171	AA523503	dbEST	Sequence 7222	AA643685	dbEST
Sequence 7172	AA525960	dbEST	Sequence 7223	AA644700	dbEST
Sequence 7173	AA526187	dbEST	Sequence 7224	AA659764	dbEST
Sequence 7174	AA526243	dbEST	Sequence 7225	AA662406	dbEST
Sequence 7175	AA526893	dbEST	Sequence 7226	AA678414	dbEST
Sequence 7176	AA531582	dbEST	Sequence 7227	AA683075	dbEST
Sequence 7177	AA533372	dbEST	Sequence 7228	AA687675	dbEST
Sequence 7178	AA533386	dbEST	Sequence 7229	AA689392	dbEST
Sequence 7179	AA534827	dbEST	Sequence 7230	AA694271	dbEST
Sequence 7180	AA535755	dbEST	Sequence 7231	AA699674	dbEST
Sequence 7181	AA548472	dbEST	Sequence 7232	AA700599	dbEST
Sequence 7182	AA554018	dbEST	Sequence 7233	AA701980	dbEST
Sequence 7183	AA555194	dbEST	Sequence 7234	AA706018	dbEST
Sequence 7184	AA557408	dbEST	Sequence 7235	AA706860	dbEST
Sequence 7185	AA564238	dbEST	Sequence 7236	AA715000	dbEST
Sequence 7186	AA565272	dbEST	Sequence 7237	AA719797	dbEST
Sequence 7187	AA568518	dbEST	Sequence 7238	AA723045	dbEST
Sequence 7188	AA568538	dbEST	Sequence 7239	AA723521	dbEST
Sequence 7189	AA569493	dbEST	Sequence 7240	AA723584	dbEST
Sequence 7190	AA574227	dbEST	Sequence 7241	AA731872	dbEST
Sequence 7191	AA577490	dbEST	Sequence 7242	AA737421	dbEST
Sequence 7192	AA579160	dbEST	Sequence 7243	AA737949	dbEST
Sequence 7193	AA579617	dbEST	Sequence 7244	AA740610	dbEST
Sequence 7194	AA581892	dbEST	Sequence 7245	AA746487	dbEST
Sequence 7195	AA582930	dbEST	Sequence 7246	AA746824	dbEST
Sequence 7196	AA584026	dbEST	Sequence 7247	AA747091	dbEST
Sequence 7197	AA585278	dbEST	Sequence 7248	AA767779	dbEST
Sequence 7198	AA585430	dbEST	Sequence 7249	AA768109	dbEST
Sequence 7199	AA586522	dbEST	Sequence 7250	AA769865	dbEST
Sequence 7200	AA587348	dbEST	Sequence 7251	AA7792 <b>78</b>	dbEST
Sequence 7201	AA592908	dbEST	Sequence 7252	AA780301	dbEST
Sequence 7202	AA594976	dbEST	Sequence 7253	AA782769	dbEST
Sequence 7203	AA595339	dbEST	Sequence 7254	AA804740	dbEST
Sequence 7204	AA595581	dbEST	Sequence 7255	AA805604	dbEST
Sequence 7205	AA599534	dbEST	Sequence 7256	AA808349	dbEST
Sequence 7206	AA599711	dbEST	Sequence 7257	AA808582	dbEST
Sequence 7207	AA600060	dbEST	Sequence 7258	AA813708	dbEST
Sequence 7208	AA602634	dbEST	Sequence 7259	AA825194	dbEST
Sequence 7209	AA602758	db <b>EST</b>	Sequence 7260	AA827738	dbEST
Sequence 7210	AA604815	dbEST	Sequence 7261	AA827816	dbEST
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Sequence 7262	AA829850	dbEST	Sequence 7313	A1074107	dbEST
Sequence 7263	AA829973	dbEST	Sequence 7314	AI076593	dbEST
Sequence 7264	AA830635	dbEST	Sequence 7315	AI077317	dbEST
Sequence 7265	AA835708	dbEST	Sequence 7316	AI081842	dbEST
Sequence 7266	AA836233	dbEST	Sequence 7317	AI081918	dbEST
Sequence 7267	AA836306	dbEST	Sequence 7318	AI085195	dbEST
Sequence 7268	AA836962	dbEST	Sequence 7319	AI087077	dbEST
Sequence 7269	AA837401	dbEST	Sequence 7320	AI087287	dbEST
Sequence 7270	AA845575	dbEST	Sequence 7321	AI091935	dbEST
Sequence 7271	AA846308	dbEST	Sequence 7322	AI093409	dbEST
Sequence 7272	AA846626	dbEST	Sequence 7323	Al097545	dbEST
Sequence 7273	AA847348	dbEST	Sequence 7324	Al114631	dbEST
Sequence 7274	AA852908	dbEST	Sequence 7325	AI123229	dbEST
Sequence 7275	AA864406	dbEST	Sequence 7326	Al125607	dbEST
Sequence 7276	AA865913	dbEST	Sequence 7327	Al131097	dbEST
Sequence 7277	AA866035	dbEST	Sequence 7328	Al 138629	dbEST
Sequence 7278	AA872969	dbEST	Sequence 7329	Al139904	dbEST
Sequence 7279	AA876384	dbEST	Sequence 7330	Al144216	dbEST
Sequence 7280	AA876730	dbEST	Sequence 7331	Al 147191	dbEST
Sequence 7281	AA886406	dbEST	Sequence 7332	Al 174867	dbEST
Sequence 7282	AA887224	dbEST	Sequence 7333	Al 183897	dbEST
Sequence 7283	AA894408	dbEST	Sequence 7334	Al184113	dbEST
Sequence 7284	AA912233	dbEST	Sequence 7335	Al208430	dbEST
Sequence 7285	AA916635	dbEST	Sequence 7336	Al214050	dbEST
Sequence 7286	AA936792	dbEST	Sequence 7337	Al217019	dbEST
Sequence 7287	AA960820	dbEST	Sequence 7338	Al217557	dbEST
Sequence 7288	AA974826	dbEST	Sequence 7339	Al222506	dbEST
Sequence 7289	AA994034	dbEST	Sequence 7340	AI240543	dbEST
Sequence 7290	AA994694	dbEST	Sequence 7341	Al241893	dbEST
Sequence 7291	Al002321	dbEST	Sequence 7342	Al246173	dbEST
Sequence 7292	AI016106	dbEST	Sequence 7343	Al249257	dbEST
Sequence 7293	Al017779	dbEST	Sequence 7344	Al249877	dbEST
Sequence 7294	AI017809	dbEST	Sequence 7345	Al252562	dbEST
Sequence 7295	AI018161	dbEST	Sequence 7346	Al262104	dbEST
Sequence 7296	AI018686	dbEST	Sequence 7347	Al262460	dbEST
Sequence 7297	Al028107	dbEST	Sequence 7348	Al263731	dbEST
Sequence 7298	AI032779	dbEST	Sequence 7349	Al266643	dbEST
Sequence 7299	AI033863	dbEST	Sequence 7350	Al271269	dbEST
Sequence 7300	AI039433	dbEST	Sequence 7351	Al275175	dbEST
Sequence 7301	AI039804	dbEST	Sequence 7352	Al280561	dbEST
Sequence 7302	AI041630	dbEST	Sequence 7353	Al281777	dbEST
Sequence 7303	AI042066	dbEST	Sequence 7354	AI283385	dbEST
Sequence 7304	Al042354	dbEST	Sequence 7355	Al301933	dbEST
Sequence 7305	AI051210	dbEST	Sequence 7356	Al302526	dbEST
Sequence 7306	AI051983	dbEST	Sequence 7357	AJ305530	dbEST
Sequence 7307	AI051994	dbEST	Sequence 7358	Al306421	dbEST
Sequence 7308	Al053744	dbEST	Sequence 7359	Al309025	dbEST
Sequence 7309	Al054344	dbEST	Sequence 7360	AJ309843	dbEST
Sequence 7310	AI056927	dbEST	Sequence 7361	Al310484	dbEST
Sequence 7311	AI057585	dbEST	Sequence 7362	Al338222	dbEST
Sequence 7312	Al066753	dbEST	Sequence 7363	Al340533	dbEST
					:

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Sequence 7364	AI340627	dbEST	Sequence 7415	AI630345	dbEST
Sequence 7365	Al345347	dbEST	Sequence 7416	Al630891	dbEST
Sequence 7366	Al345370	dbEST	Sequence 7417	A1635478	dbEST
Sequence 7367	Al345478	dbEST	Sequence 7418	A1636626	dbEST
Sequence 7368	Al347461	dbEST	Sequence 7419	AI637584	dbEST
Sequence 7369	Al349256	dbEST	Sequence 7420	Al653541	dbEST
Sequence 7370	Al357779	dbEST	Sequence 7421	A1656402	dbEST
Sequence 7371	AI360123	dbEST	Sequence 7422	AI659386	dbEST
Sequence 7372	Al363118	dbEST	Sequence 7423	A1670009	dbEST
Sequence 7373	AI364592	dbEST	Sequence 7424	Al671642	dbEST
Sequence 7374	AI367730	dbEST	Sequence 7425	AI687362	dbEST
Sequence 7375	AI373565	dbEST	Sequence 7426	AI696693	dbEST
Sequence 7376	AI375751	dbEST	Sequence 7427	AI698092	dbEST
Sequence 7377	A1394420	dbEST	Sequence 7428	A1720132	dbEST
Sequence 7378	Al394469	dbEST	Sequence 7429	A1741446	dbEST
Sequence 7379	AI400282	dbEST	Sequence 7430	AI741861	dbEST
Sequence 7380	Al401513	dbEST	Sequence 7431	AI745218	dbEST
Sequence 7381	Al420373	dbEST	Sequence 7432	A1753459	dbEST
Sequence 7382	Al421290	dbEST	Sequence 7433	AI796562	dbEST
Sequence 7383	Al421401	dbEST	Sequence 7434	A1799699	dbEST
Sequence 7384	Al431327	dbEST	Sequence 7435	AI801589	dbEST
Sequence 7385	Al431923	dbEST	Sequence 7436	A1802542	dbEST
Sequence 7386	Al434109	dbEST	Sequence 7437	AI811144	dbEST
Sequence 7387	A1434949	dbEST	Sequence 7438	A1826427	dbEST
Sequence 7388	Al439452	dbEST	Sequence 7439	Al827129	dbEST
Sequence 7389	A1440238	dbEST	Sequence 7440	AI827550	dbEST
Sequence 7390	AI446003	dbEST	Sequence 7441	A1832609	dbEST
Sequence 7391	Al446426	dbEST	Sequence 7442	AI866551	dbEST
Sequence 7392	Al453477	dbEST	Sequence 7443	A1913323	dbEST
Sequence 7393	AI457107	dbEST	Sequence 7444	Al916374	dbEST
Sequence 7394	AI457410	dbEST	Sequence 7445	Al916419	dbEST
Sequence 7395	A1458799	dbEST	Sequence 7446	Al921254	dbEST
Sequence 7396	Al472566	dbEST	Sequence 7447	Al922114	dbEST
Sequence 7397	Al476075	dbEST	Sequence 7448	Al923485	dbEST
Sequence 7398	Al492953	dbEST	Sequence 7449	A1928889	dbEST
Sequence 7399	AI497636	dbEST	Sequence 7450	Al932951	dbEST
Sequence 7400	A1499584	dbEST	Sequence 7451	Al937150	dbEST
Sequence 7401	AI523964	dbEST	Sequence 7452	AI954406	dbEST
Sequence 7402	A1524304	dbEST	Sequence 7453	Al961592	dbEST
Sequence 7403	AI537244	dbEST	Sequence 7454	AI968311	dbEST
Sequence 7404	AI538342	dbEST	Sequence 7455	AL036165	dbEST
Sequence 7405	AI557082	dbEST	Sequence 7456	AL036575	dbEST
Sequence 7406	A1560184	dbEST	Sequence 7457	AL037646	dbEST
Sequence 7407	A1560756	dbEST	Sequence 7458	AL043071	dbEST
Sequence 7408	AI570521	dbEST	Sequence 7459	AL045040	dbEST
Sequence 7409	AI572096	dbEST	Sequence 7460	AL045500	dbEST
Sequence 7410	AI573218	dbEST	Sequence 7461	AL046648	dbEST
Sequence 7411	AI582912	dbEST	Sequence 7462	AL110384	dbEST
Sequence 7412	AI583257	dbEST	Sequence 7463	AL119319	dbEST
Sequence 7413	Al611738	dbEST	Sequence 7464	AW009700	dbEST
Sequence 7414	Al625204	dbEST	Sequence 7465	AW009886	dbEST

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#### **Table 3-1**

Sequence 7466	AW015094	dbEST
Sequence 7467	AW016443	dbEST
Sequence 7468	AW016546	dbEST
Sequence 7469	AW020245	dbEST
Sequence 7470	AW021402	dbEST
Sequence 7471	AW023884	dbEST
Sequence 7472	AW025130	dbEST
Sequence 7473	AW043908	dbEST
Sequence 7474	AW075324	dbEST
Sequence 7475	AW079132	dbEST
Sequence 7476	AW080456	dbEST
Sequence 7477	AW084350	dbEST
Sequence 7478	AW084663	dbEST
Sequence 7479	AW117855	dbEST
Sequence 7480	AW130776	dbEST
Sequence 7481	AW149504	dbEST
Sequence 7482	AW161853	dbEST
Sequence 7483	AW167569	dbEST
Sequence 7484	AW169156	dbEST
Sequence 7485	AW183848	dbEST
Sequence 7486	D52802	dbEST
Sequence 7487	D55565	dbEST
Sequence 7488	F28764	dbEST
Sequence 7489	H47715	dbEST
Sequence 7490	H48257	dbEST
Sequence 7491	H69070	dbEST
Sequence 7492	H93650	dbEST
Sequence 7493	N30570	dbEST
Sequence 7494	N31301	dbEST
Sequence 7495	N75979	dbEST
Sequence 7496	R48046	dbEST
Sequence 7497	W05661	dbEST
Sequence 7498	W35236	dbEST
Sequence 7499	W37602	dbEST
Sequence 7500	W73817	dbEST
Sequence 7501	W79668	dbEST
Sequence 7502	W79717	dbEST
Sequence 7503	Q65520	NUCPATENT
Sequence 7504	V05728	NUCPATENT
Sequence 7505	V63195	NUCPATENT
Sequence 7506	X00656	NUCPATENT
Sequence 7507	X08687	NUCPATENT
Sequence 7508	X24921	NUCPATENT
Sequence 7509	X27427	NUCPATENT
Sequence 7510	X85038	NUCPATENT
Sequence 7511	X89652	NUCPATENT

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#### Table 3-2

Sequence 1240

Sequence 2802

CCGCGGTGGCGGCCGAGGTNCCAANGTCAAGTCACCATAATGTNTTTNACCTGTTTCACA ATGTTCTACATTATCGTGGAAATGTTGACAAGGTCTCNAGCAAAGACTGGATCTNCANTG GNTTTAAAATACAATNNTCCATTTCGGAAATTGANGGAAAAAGATCTTTNNCAANCTGGA AGAGTCTTCTTGTAAATAGGAATCACNGGGCTACACCTGCTGACGACCTTGGGTTTN

Sequence 2803

TTAGGGCGAATTGGAGCTCCCCGCGGTGGCGGCCGAGGTACTATATGGACACCAACATGG AGCANAAATGAGGGCAGCAGTGTCCAATCTCATTCCAGGTGAGAAGTTGCACAGTGTCC AATAGGTGCATACATCTCCTTAGTAAGTAGTTGTGATTAACAATGAAATAGAAATGAAAA ATATATTTTTTATTTATGTGTATTATATTTTTCAAGCAGCTAATAAGTTGGTAGGACAT AATATTTAATTCGTTGGGGACCTAATTATTTATAAATTGAATGGTTAGATATTTTTTG GCCTAAGCCACCATGAAAAAATCACTGAGGCGCTAAGGGAAACATGAACTAAGAAGCCCT TCTGAGTCTCTGTTTTCTCAACTGTAAGATGAANAAACCTGCTCCACCTTCCTCTTGCGA TTGGTGAACGGCAGATTTGAGATACTGCGCAGAATGGACCTTAT

Sequence 2804

Sequence 2805

CTATAGGGCGAATTGGAGCTCCCCGCGGTGGCGGCCCCCGGGCATGGTACGCGGGGAGG TGTTGTTCCGTGAATCAGCACATCAATTGCAGCATTGTGGCTACCAGGGGGTCAGGATGC GGNCGGTGGAGCCCTCTGGCCTTTGTGTGGTAGCCGAGGACTCTGTGTCAGCTGACCGTT TTCCGGNAAACTTTCGTGCGAGACTCACATCTTGGAAATTCAAATACTCAATANCTCTCG AATTCTAGGAATCTTGAGAAGAGCCCTGGATTAANGATTCANACATGGGCCCTNANATNG NTATGGCATTGCTGGTTCTACCAACGTGACAG

Sequence 4710

CCGGGCAGGTACCAGAACTTCAGCAAAGGAAGCAGACACAGGTCGGAGGAAGCCTACATA
GACCCCATTGCCATGGAGTATTACAACTGGGGGGCGGTTCTCGAAGCCCCCAGAAGATGAT
GATGCCAATTCCTACGAGAATGTGCTCATTTGCAAGCAGAAAACCACAGAGACAGGTGCC
CAGCAGGAGGGCATAGGTGGCCTCTGCAGAGGGGACCTCAGCCTGTCACTGGCCCTGAAG
ACTGGCCCCACTTCTGGTCTCTGTCCCCCTGCCTCCCGGAAGAAGATGAGGAATCTGAG
GATTATCAGAACTCAGCATCCATCCATCAGTGGCACGAGTCCAGGAAG

Sequence 4711

Sequence 4712

CGCTCCGCCCGCTNACACGGCTTCCTGGGCGGTCCCTCCACCTGTTGCTTCAGGTCCT GCAAGCCCTTGCTTGCCATGGCTTCGGGGTATCTGTGGAGTCNTCAAGANCANCTGGAGC CACNTTGGATCCTNCCCAAAGTGGCCCCCGCGTACCTTGGCCCGTNTAAAACTATGTGGA ATCCCCCGGGCCTGCANGAATTCCATATCAAAGCTTATNGATACCCGGCGACCCTCGAGG GGG

Sequence 4713

CCGCGGTGGCGGCCGAGGTACGCCGGGTCATTACTACAGGAAAAACTGTTCTCTTCTGTG
GCACAGAGAACCCTGCTTCAAAGCAGAAGTAGCAGTTCCGGAGTCCAGCTGGCTAAAACT
CATCCCAGAGGATAATGGCAACCCATGCCTTAGAAATCGCTGGGCTGTTTCTTGGTGGT
TTGGAATGGTGGGCACAGTGGCTGTCACTGTCATGCCTCAGTGGAGAGTGTCGGCCTTCA
TTGAAAACAACATCGTGGTTTTTGAAAACTTCTGGGAAGGACTGTGGATGAATTGCGTGA
GGCAGGCTAACATCAGGATGCAGTTGCAAAATCTATGATTCCCTGCTGGCTCTTTCTCCG
GACCTACAGGCAGCCAGAGGACTGATGTTGCGCTGCTTCCTTGGCTTT
CATGATGGC

Sequence 4714

CGCTGATGTTGGCCANGCGAGAGGGCCACGTTCTCCTGAGTGTTCAGAAGGGTCTTACAG
ACGTCCAGGGGGGTCGTGGCGGCCGGCGAGGGCCCCGGCCAGCCCGCCTGAGATGATG
TGGGACTGCGGGTTGTAGGTCCGGTGGGGGTTGACCTGCTCCTGCAGGAACTCATAGGTG
ATGAAGTGGATGGACTGGAAGGGGATGTTCATGGTCAGCTGCGTGGTGTAGCTCCGGTAG
AAGGCCCCCAACCCCTNGGTCCTCCACACCGTCCGGATGCAACTGATTGCTGACCGGTGC
TGCNAGTTGTACCTGCCCGGGCGGCCGCTCTAGAACTA

Sequence 4715

GCCGCCGGCCAGGTACAAACTTAGAAGAAAATNGGAAGATAGAAACAAGATAGAAAATGA
AAATATTGTCAAGAGTTCAGATAGAAAATGAAAAACAAGCTAAGACAAGTATTGGAGAA
GTATAGAAGATAGAAAAATTAGAGCCAAAAATTGGATAAAATAGACACTGAAAAAATGAG
GAAATTATTGGTAACCAATTTATTTTAAAAGCCCATCAATTTAATTTCTGGTGGTGCAGA
AGTTAGAAGGTAAAGCTTGAGAAGATGAGGGTGTTTACCGTAGACCNGAACCAATTTAGA
AGAATACTTGAAGCTAGAAGGGGAAGTTGGTTAAAAATCACATCAAAAAGCTACTAAAAG
GACTGGTGTAATTTTAAAAAAAAACTAAGGCAGAAGGCTTTTGGAAGAAGTTAGAAGAATT
TGGAAGGCCTTAAATATAGTAGCTTAATTTGAAAAAATGTGAAGGACCTTTCGTAACCGG
AAGGTNATTTCAAGATNAAAGAAGTAATTTACCAACCTTAATGGTTTTTGCCTTTTGGGC
TTTTGAGTTTAAAGAATTANT

Sequence 4716

Sequence 4717

Sequence 4718

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TGGCAACAGAGATACCCCACTTCACTGGGTAGCCTTTAAGAACAATGCTGAGTGTGTGCGGGCTCTCCTAGAGAGCGGGGCCTCTGTCAATGCCCTGGA

Sequence 4719

Sequence 4720

Sequence 4721

Sequence 4722

GGTGGCGGCCGAGGTACCTCAAAAGTGGGGAAATCCCCATATAACTGANGACGAAGGCAG TTCAGAAGTTCATTGATTTGCCCTAAGGTTCCTCAATTTGCAAACGTCAGGCCAATGATC CAACCCCAGGTATGTTTGGCAGTGAAGGACCAGTTGAGTCATAGCTGCAAGTAACCACCC TGCAGTGGTCCCTATCTTGGCCGTTAGCTTACATTGACATTTAACACTCAAATTTACTCA GTAACACCAGCTATCATGTTTTCCACTAAAACTCCACAGCATTCTGGCAACTTTCTATT TTAGAGCAATAAAGTAAATTGTTAGCATCCCTTTGACATATAAATATTTCTACAAATAGT AATTCTCTAGCCATTCATTTGGAGTATTTAAAACTCAACATTCATAGCACATTTTATGGT GACAAAGAACTTATGTTCAGAACACAAAAAATAAGTCGTACCTGCCCGGGCGGCCGGTCT

Sequence 4723

ACAAAACCAATCTACCTGATGAAAACTCCGTTCCCTTCTCGCCAGAAACATAAAATGCG ATGGAGCTACGGCCACCGCTGCCGAGACAAAATGGCGCCGACCCCGCGTACCT

Sequence 6889

Sequence 6890

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ATGGCCTACCCAACATCCATTCTCTACTCCCTCTACTCTGATGGCACCCG

Sequence 6891

Sequence 6892

CTCCCCGCGTGGCGGCCGAGGTACGCGGGGCTCTATATAAGTGGGCAGNGGCCGNGACT GCGCCCAGACACTGACCTTCAGCGCCCTNGGCTCCAGCGCCATGGCGCCCTCCAGGAAGTT CTTCGTTGGGGGAAACTGGAAGATGAACGGACGGAAGCAGANTCTGGGGGAGCTCATCGG CACTCTNAACGCGGCCAA

Sequence 6893

Sequence 6894

Sequence 6895

Sequence 6896

CTACTTAGGGCGAATTGGAGCTCCCCGCGGTGGCGGCCCCCGGGCAGGTACTCTGGTCA
TCTTCGTTCGTTTGGTCGTAAGGTGTTAACTATTTTCACTTCCCATATCACAAAGTTA
GTCCACAGGAGGAGCTGGTGGATCTTGTCCATTATGAGGACTGGTTGGCTTTCCAGGTAG
GCTGGATCCTNTTAGATTAGGAGGGTCTCAGTAAGAACAGGATCAATGCAATAACCATCC
AGGCTACCAAGATNATTGTAACACTGATGCCNTTATNACCAGAGGGTCCCGGTAATTNCT
GAANACACTCTGTGTCTGTGCAGTATGATTGGGACTGCCGTANCANATTGNTCAGTCTTC
TCANTGCATNGTTTCATGAGAGCAAAACACATTNACAGGCATCAAATTCCACCTTCTTGC
CCATTATTACCCAGCTTGGATTTTTACTTGGACCTNTTTTTAAAATCCCNCC

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Table 3-3

Sequence #8         Acc No         Database hit         GI Number           Sequence 7512         A14133         Genbank         490127           Sequence 7514         A26895         Genbank         1247450           Sequence 7516         A29216         Genbank         1247460           Sequence 7516         A33292         Genbank         1247528           Sequence 7517         A36460         Genbank         2293778           Sequence 7518         A43521         Genbank         2298710           Sequence 7520         A80239         Genbank         3713993           Sequence 7520         A80239         Genbank         6092971           Sequence 7521         A81324         Genbank         6092971           Sequence 7522         A98295         Genbank         6731650           Sequence 7524         AB000360         Genbank         2547041           Sequence 7525         AB000516         Genbank         2723379           Sequence 7525         AB000584         Genbank         273379           Sequence 7527         AB000624         Genbank         2467299           Sequence 7530         AB001601         Genbank         2467299           Sequence 7531	Comun	A . N	D 4 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	O. N.
Sequence 7513         A26481         Genbank         1247452           Sequence 7514         A26595         Genbank         1247460           Sequence 7516         A32921         Genbank         125728           Suquence 7516         A33292         Genbank         1567852           Sequence 7517         A36460         Genbank         2298778           Sequence 7518         A43521         Genbank         2298710           Sequence 7520         A80239         Genbank         6092971           Sequence 7521         A81324         Genbank         6731650           Sequence 7522         A98295         Genbank         6731650           Sequence 7522         A8000360         Genbank         2547041           Sequence 7524         AB000468         Genbank         2547041           Sequence 7525         AB000584         Genbank         2723379           Sequence 7527         AB000624         Genbank         2743379           Sequence 7529         AB001623         Genbank         2467299           Sequence 7529         AB001623         Genbank         224600           Sequence 7531         AB002293         Genbank         2224530           Sequence 7534				
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Sequence 7543         AB002365         Genbank         2224674           Sequence 7544         AB002368         Genbank         2224680           Sequence 7545         AB002374         Genbank         228 3484           Sequence 7546         AB002375         Genbank         226 3486           Sequence 7547         AB002377         Genbank         226 3486           Sequence 7548         AB002377         Genbank         2224706           Sequence 7549         AB002382         Genbank         2224708           Sequence 7550         AB002387         Genbank         2224718           Sequence 7551         AB002387         Genbank         2224718           Sequence 7551         AB002387         Genbank         2081606           Sequence 7552         AB003476         Genbank         2081606           Sequence 7553         AB003730         Genbank         2627128           Sequence 7555         AB006630         Genbank         2564331           Sequence 7556         AB006630         Genbank         2564331           Sequence 7557         AB007447         Genbank         2463530           Sequence 7558         AB007510         Genbank         2463576           Sequ				
Sequence 7544         AB002368         Genbank         2224680           Sequence 7545         AB002374         Genbank         228 484           Sequence 7546         AB002375         Genbank         226 486           Sequence 7547         AB002377         Genbank         6634024           Sequence 7548         AB002381         Genbank         2224706           Sequence 7549         AB002382         Genbank         2224708           Sequence 7550         AB002387         Genbank         2224718           Sequence 7551         AB002559         Genbank         2224718           Sequence 7552         AB003476         Genbank         2081606           Sequence 7553         AB003476         Genbank         2627128           Sequence 7554         AB005754         Genbank         2627128           Sequence 7554         AB006630         Genbank         2564331           Sequence 7556         AB006651         Genbank         2564331           Sequence 7557         AB007447         Genbank         2463530           Sequence 7558         AB007510         Genbank         2463576           Sequence 7550         AB007889         Genbank         2662150           Sequenc				
Sequence 7545         AB002374         Genbank         228 484           Sequence 7546         AB002375         Genbank         228 486           Sequence 7547         AB002377         Genbank         6634024           Sequence 7548         AB002381         Genbank         2224706           Sequence 7549         AB002382         Genbank         2224708           Sequence 7550         AB002387         Genbank         2224718           Sequence 7551         AB002559         Genbank         2081606           Sequence 7552         AB003476         Genbank         2081606           Sequence 7553         AB003730         Genbank         2627128           Sequence 7554         AB005754         Genbank         6456482           Sequence 7554         AB006630         Genbank         2564331           Sequence 7556         AB006651         Genbank         2463530           Sequence 7558         AB007510         Genbank         2463576           Sequence 7559         AB007859         Genbank         2463576           Sequence 7560         AB007882         Genbank         2887418           Sequence 7561         AB007895         Genbank         2662150           Sequenc				
Sequence 7546         AB002375         Genbank         2253486           Sequence 7547         AB002377         Genbank         6634024           Sequence 7548         AB002381         Genbank         2224706           Sequence 7549         AB002382         Genbank         2224708           Sequence 7550         AB002387         Genbank         2224718           Sequence 7551         AB002559         Genbank         1944129           Sequence 7552         AB003476         Genbank         2081606           Sequence 7553         AB003730         Genbank         2627128           Sequence 7554         AB005754         Genbank         2564331           Sequence 7555         AB006630         Genbank         2564331           Sequence 7556         AB006651         Genbank         2463530           Sequence 7557         AB007447         Genbank         2463530           Sequence 7558         AB007510         Genbank         2463576           Sequence 7560         AB007882         Genbank         2887418           Sequence 7561         AB007895         Genbank         2662150           Sequence 7563         AB007990         Genbank         2662156           Sequenc				
Sequence 7547         AB002377         Genbank         6634024           Sequence 7548         AB002381         Genbank         2224706           Sequence 7549         AB002382         Genbank         2224708           Sequence 7550         AB002387         Genbank         2224718           Sequence 7551         AB002559         Genbank         1944129           Sequence 7552         AB003476         Genbank         2081606           Sequence 7553         AB003730         Genbank         2627128           Sequence 7554         AB005754         Genbank         6456482           Sequence 7555         AB006630         Genbank         2564331           Sequence 7556         AB006651         Genbank         3201570           Sequence 7557         AB007447         Genbank         2463530           Sequence 7558         AB007510         Genbank         2463576           Sequence 7550         AB007859         Genbank         2887418           Sequence 7561         AB007895         Genbank         2662150           Sequence 7562         AE007899         Genbank         2662158           Sequence 7564         AB007900         Genbank         2662160           Sequenc				
Sequence 7548         AB002381         Genbank         2224706           Sequence 7549         AB002382         Genbank         2224708           Sequence 7550         AB002387         Genbank         2224718           Sequence 7551         AB002559         Genbank         1944129           Sequence 7552         AB003476         Genbank         2081606           Sequence 7553         AB003730         Genbank         2627128           Sequence 7554         AB005754         Genbank         6456482           Sequence 7555         AB006630         Genbank         2564331           Sequence 7556         AB006651         Genbank         3201570           Sequence 7557         AB007447         Genbank         2463530           Sequence 7558         AB007510         Genbank         2463576           Sequence 7559         AB007859         Genbank         6634028           Sequence 7561         AB007882         Genbank         2887418           Sequence 7562         AE007899         Genbank         2662150           Sequence 7563         AB007900         Genbank         2662160           Sequence 7564         AB007920         Genbank         3413863				
Sequence 7549         AB002382         Genbank         2224708           Sequence 7550         AB002387         Genbank         2224718           Sequence 7551         AB002559         Genbank         1944129           Sequence 7552         AB003476         Genbank         2081606           Sequence 7553         AB003730         Genbank         2627128           Sequence 7554         AB005754         Genbank         6456482           Sequence 7555         AB006630         Genbank         2564331           Sequence 7556         AB006651         Genbank         3201570           Sequence 7557         AB007447         Genbank         2463530           Sequence 7558         AB007510         Genbank         2463576           Sequence 7559         AB007859         Genbank         6634028           Sequence 7561         AB007882         Genbank         2887418           Sequence 7562         AE007899         Genbank         2662150           Sequence 7563         AB007900         Genbank         2662160           Sequence 7564         AB007920         Genbank         3413863				
Sequence 7550         AB002387         Genbank         2224718           Sequence 7551         AB002559         Genbank         1944129           Sequence 7552         AB003476         Genbank         2081606           Sequence 7553         AB003730         Genbank         2627128           Sequence 7554         AB005754         Genbank         6456482           Sequence 7555         AB006630         Genbank         2564331           Sequence 7556         AB006651         Genbank         3201570           Sequence 7557         AB007447         Genbank         2463530           Sequence 7558         AB007510         Genbank         2463576           Sequence 7559         AB007859         Genbank         6634028           Sequence 7560         AB007882         Genbank         2887418           Sequence 7561         AB007895         Genbank         2662150           Sequence 7562         AE007899         Genbank         2662158           Sequence 7564         AB007900         Genbank         2662160           Sequence 7564         AB007920         Genbank         3413863			_	
Sequence 7551         AB002559         Genbank         1944129           Sequence 7552         AB003476         Genbank         2081606           Sequence 7553         AB003730         Genbank         2627128           Sequence 7554         AB005754         Genbank         6456482           Sequence 7555         AB006630         Genbank         2564331           Sequence 7556         AB006651         Genbank         3201570           Sequence 7557         AB007447         Genbank         2463530           Sequence 7558         AB007510         Genbank         2463576           Sequence 7559         AB007859         Genbank         6634028           Sequence 7560         AB007882         Genbank         2887418           Sequence 7561         AB007895         Genbank         2662150           Sequence 7562         AE007899         Genbank         2662158           Sequence 7564         AB007900         Genbank         2662160           Sequence 7564         AB007920         Genbank         3413863				
Sequence 7552         AB003476         Genbank         2081606           Sequence 7553         AB003730         Genbank         2627128           Sequence 7554         AB005754         Genbank         6456482           Sequence 7555         AB006630         Genbank         2564331           Sequence 7556         AB006651         Genbank         3201570           Sequence 7557         AB007447         Genbank         2463530           Sequence 7558         AB007510         Genbank         2463576           Sequence 7559         AB007859         Genbank         6634028           Sequence 7560         AB007882         Genbank         2887418           Sequence 7561         AB007895         Genbank         2662150           Sequence 7562         AE007899         Genbank         2662158           Sequence 7563         AB007900         Genbank         2662160           Sequence 7564         AB007920         Genbank         3413863				
Sequence 7553         AB003730         Genbank         2627128           Sequence 7554         AB005754         Genbank         6456482           Sequence 7555         AB006630         Genbank         2564331           Sequence 7556         AB006651         Genbank         3201570           Sequence 7557         AB007447         Genbank         2463530           Sequence 7558         AB007510         Genbank         2463576           Sequence 7559         AB007859         Genbank         6634028           Sequence 7560         AB007882         Genbank         2887418           Sequence 7561         AB007895         Genbank         2662150           Sequence 7562         AE007899         Genbank         2662158           Sequence 7563         AB007900         Genbank         2662160           Sequence 7564         AB007920         Genbank         3413863				
Sequence 7554         AB005754         Genbank         6456482           Sequence 7555         AB006630         Genbank         2564331           Sequence 7556         AB006651         Genbank         3201570           Sequence 7557         AB007447         Genbank         2463530           Sequence 7558         AB007510         Genbank         2463576           Sequence 7559         AB007859         Genbank         6634028           Sequence 7560         AB007882         Genbank         2887418           Sequence 7561         AB007895         Genbank         2662150           Sequence 7562         AE007899         Genbank         2662158           Sequence 7563         AB007900         Genbank         2662160           Sequence 7564         AB007920         Genbank         3413863				
Sequence 7555         AB006630         Genbank         2564331           Sequence 7556         AB006651         Genbank         3201570           Sequence 7557         AB007447         Genbank         2463530           Sequence 7558         AB007510         Genbank         2463576           Sequence 7559         AB007859         Genbank         6634028           Sequence 7560         AB007882         Genbank         2887418           Sequence 7561         AB007895         Genbank         2662150           Sequence 7562         AE007899         Genbank         2662158           Sequence 7563         AB007900         Genbank         2662160           Sequence 7564         AB007920         Genbank         3413863	Sequence 7554			
Sequence 7556         AB006651         Genbank         3201570           Sequence 7557         AB007447         Genbank         2463530           Sequence 7558         AB007510         Genbank         2463576           Sequence 7559         AB007859         Genbank         6634028           Sequence 7560         AB007882         Genbank         2887418           Sequence 7561         AB007895         Genbank         2662150           Sequence 7562         AE007899         Genbank         2662158           Sequence 7563         AB007900         Genbank         2662160           Sequence 7564         AB007920         Genbank         3413863				
Sequence 7557         AB007447         Genbank         2463530           Sequence 7558         AB007510         Genbank         2463576           Sequence 7559         AB007859         Genbank         6634028           Sequence 7560         AB007882         Genbank         2887418           Sequence 7561         AB007895         Genbank         2662150           Sequence 7562         AE007899         Genbank         2662158           Sequence 7563         AB007900         Genbank         2662160           Sequence 7564         AB007920         Genbank         3413863			_	
Sequence 7558         AB007510         Genbank         2463576           Sequence 7559         AB007859         Genbank         6634028           Sequence 7560         AB007882         Genbank         2887418           Sequence 7561         AB007895         Genbank         2662150           Sequence 7562         AE007899         Genbank         2662158           Sequence 7563         AB007900         Genbank         2662160           Sequence 7564         AB007920         Genbank         3413863			_	
Sequence 7559         AB007859         Genbank         6634028           Sequence 7560         AB007882         Genbank         2887418           Sequence 7561         AB007895         Genbank         2662150           Sequence 7562         AE007899         Genbank         2662158           Sequence 7563         AB007900         Genbank         2662160           Sequence 7564         AB007920         Genbank         3413863	•			
Sequence 7560         AB007882         Genbank         2887418           Sequence 7561         AB007895         Genbank         2662150           Sequence 7562         AE007899         Genbank         2662158           Sequence 7563         AB007900         Genbank         2662160           Sequence 7564         AB007920         Genbank         3413863			_	
Sequence 7561         AB007895         Genbank         2662150           Sequence 7562         AE 007899         Genbank         2662158           Sequence 7563         AB007900         Genbank         2662160           Sequence 7564         AB007920         Genbank         3413863				
Sequence 7562         A=007899         Genbank         2662158           Sequence 7563         AB007900         Genbank         2662160           Sequence 7564         AB007920         Genbank         3413863				
Sequence 7563         AB007900         Genbank         2662160           Sequence 7564         AB007920         Genbank         3413863	· · · · · · · · · · · · · · · · · · ·			
Sequence 7564 AB007920 Genbank 3413863				
Sequence 7565 AB007930 Genbank 3413883			_	3413883
Sequence 7566 AB007931 Genbank 3413885				

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Sequence 7567	ABC07944	Genbank	3413911
Sequence 7568	ABC07957	Genbank	3413931
Sequence 7569	ABC07965	Genbank	3413940
Sequence 7570	AB009398	Genbank	3618342
Sequence 7571	AB011086	Genbank	3043551
Sequence 7572	AB011087	Genbank	3043553
Sequence 7573	AB011096	Genbank	3043571
Sequence 7574	AB011113	Genbank	3043605
Sequence 7575	AB011144	Genbank	3043667
Sequence 7576	AB011149	Genbank	3043677
Sequence 7577	AB011151	Genbank	3043681
Sequence 7578	AB011155	Genbank	3043689
Sequence 7579	AB011166	Genbank	3043711
Sequence 7580	AB011167	Genbank	3043713
Sequence 7581	AB011169	Genbank	3043717
Sequence 7582	AB011175	Genbank	3043729
Sequence 7583	AB011536	Genbank	3449297
Sequence 7584	AB011542	Genbank	3449309
Sequence 7585	AB012124	Genbank	4126408
Sequence 7586	AB012193	Genbank	3599675
Sequence 7387	AB012911	Genbank	3062802
Sequence 7588	AB012922	Genbank	4126426
Sequence 7589	AB014087	Genbank	5672599
Sequence 7590	AB014535	Genbank	3327083
Sequence 7591	AB014540	Genbank	3327093
Sequence 7592	AB014542	Genbank	3327097
Sequence 7593	AB014557	Genbank	3327127
Sequence 7594	AB014563	Genbank	3327139
Sequence 7595	AB014566	Genbank	3327145
Sequence 7596	AB014569	Genbank	3327151
Secuence 7597	AB014589	Genbank	3327191
Sequence 7598	AB014592	Genbank	3327197
Sequence 7599	AB014597	Genbank	3327207
Sequence 7600	AB015597	Genbank	4115713
Sequence 7601	AB017114	Genbank	4586879
Sequence 7602	AB017335	Genbank	3582440
Sequence 7603	AB017498	Genbank	3582144
Sequence 7604	AB017548	Genbank	3885363
Sequence 7605	AB018247	Genbank	4049592
Sequence 7606	AB018255	Genbank	3882144
Sequence 7607	AB018281	Genbank	3882196
Sequence 7608	AB018290	Genbank	3882214
Sequence 7609	AB018302	Genbank	3882238
Sequence 7610	AB018315	Genbank	3882264
Sequence 7611	AB018316	Genbank	3882266
Sequence 7612	AB018324	Genbank	3882282
Sequence 7613	AB018331	Genbank	3882296
Sequence 7614	AB018335	Genbank	3882304
Sequence 7615	AB018342	Genbank	3882318
Sequence 7616	AB018347	Genbank	3882328
Sequence 7617	AB018350	Genbank	3882334
Sequence 7618	AB018353	Genbank	3882340
Sequence 7619	AB018357	Genbank	6009486
Sequence 7620	AB019564	Genbank	3885367
Sequence 7821	AB019568	Genbank	3885371
Sequence 7622	AB020315	Genbank	39 <b>25828</b>

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Sequence 7623	AB020335	Genbank	6518494
Sequence 7624	AB020623	Genbank	3985929
Sequence 7625	AB020631	Genbank	4240136
Sequence 7626	AB020645	Genbank	4240164
Sequence 7627	AB020659	Genbank	4240192
	AB020662	Genbank	4240198
Sequence 7628			
Sequence 7629	AB020666	Genbank	6635134
Sequence 7630	AB020674	Genbank	4240222
Sequence 7631	AB020679	Genbank	4240232
Sequence 7632	AB020692	Genbank	4240258
Sequence 7633	AB020697	Genbank	4240268
Sequence 7634	AB020698	Genbank	4240270
Sequence 7635	AB020713	Genbank	4240300
Sequence 7636	AB020714	Genbank	4240302
Sequence 7637	AB020718	Genbank	4240310
Sequence 7638	AB020721	Genbank	4240316
Sequence 7639	AB020764	Genbank	6467144
Sequence 7640	AB020874	Genbank	4003394
Sequence 7641	AB021663	Genbank	4996450
Sequence 7041	AB022537	Genbank	6942217
Sequence 7642	AB023049	Genbank	5672604
Sequence 7643	AB023049 AB023050	Genbank	5672605
Sequence 7644			5672606
Sequence 7645	AB023051	Genbank	5672607
Sequence 7646	AB023052	Genbank	4589471
Sequence 7647	AB023137	Genbank	
Sequence 7648	AB023145	Genbank	4589487
Sequence 7649	AB023150	Genbank	4589509
Sequence 7650	AB023152	Genbank	4589513
Sequence 7651	AB023160	Genbank	4589529
Sequence 7652	AB023165	Genbank	4589539
Sequence 7653	AB023195	Genbank	4589599
Sequence 7654	AB023199	Genbank	4589607
Sequence 7655	AB023205	Genbank	4589619
Sequence 7656	AB023420	Genbank	4579908
Sequence 7657	AB024704	Genbank	4589928
Sequence 7658	AB025580	Genbank	5931568
Sequence 7659	AB026906	Genbank	5295993
Sequence 7660	AB028893	Genbank	6552364
Sequence 7661	AB028942	Genbank	5689368
Sequence 7662	AB028952	Genbank	5689394
Sequence 7663	AB028985	Genbank	5689460
Sequence 7664	AB028986	Genbank	5689462
Sequence 7665	AB028990	Genbank	5689470
Sequence 7666	AB028999	Genbank	5689488
Sequence 7667	AB029002	Genbank	5689494
Sequence 7668	AB029003	Genbank	5689496
Sequence 7669	AB029008	Genbank	5689506
Sequence 7670	AB029013	Genbank	5689516
Sequence 7671	AB029017	Genbank	5689524
Sequence 7672	AB029019	Genbank	5689528
Sequence 7673	AB029025	Genbank	5689540
Sequence 7674	AB029026	Genbank	5689542
	AB029027	Genbank	5689544
Sequence 7675	AB031051	Genbank	6683742
Sequence 7676	AB032417	Genbank	6277265
Sequence 7677		Genbank	6329896
Sequence 7678	AB032963	Genbank	0323030

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Sequence 7679	AB032975	Genbank	6330044
Sequence 7680	AB032978	Genbank	6330094
Sequence 7681	AB032983	Genbank	6330128
Sequence 7682	AB033007	Genbank	6330242
Sequence 7683	AB033011	Genbank	6330301
Sequence 7684	AB033018	Genbank	6330350
Sequence 7685	AB033023	Genbank	6330384
Sequence 7686	AB033026	Genbank	6330406
Sequence 7687	AB033027	Genbank	6330415
Sequence 7688	AB033033	Genbank	6330496
Sequence 7689	AB033051	Genbank	6330667
Sequence 7690	AB033058	Genbank	6330721
Sequence 7691	AB033070	Genbank	6330818
Sequence 7692	AB033073	Genbank	6330839
Sequence 7693	AB033078	Genbank	6330873
Sequence 7694	AB033104	Genbank	6331314
Sequence 7695	AB034205	Genbank	6899845
Sequence 7696	AB035519	Genbank	6681348
Sequence 7697	AB035698	Genbank	6970477
Sequence 7698	AB037728	Genbank	7242968
Sequence 7699	AB037743	Genbank	7243024
Sequence 7700	AB037748	Genbank	7243034
Sequence 7701	AB037748	Genbank	7243034
Sequence 7701	AB037763	Genbank	7243064
Sequence 7703	AB037775	Genbank	7243088
	AB037783	Genbank	
Sequence 7704	AB037784	Genbank Genbank	7243104
Sequence 7705			7243106
Sequence 7706	AB037797	Genbank	7243132
Sequence 7707	AB037801	Genbank	7243140
Sequence 7708	AB037807	Genbank	7243152
Sequence 7709	AB037819	Genbank	7243176
Sequence 7710	AB037844	Genbank	7243226
Sequence 7711	AB037847	Genbank	7243232
Sequence 7712	AB037856	Genbank	7243267
Sequence 7713	AC000024	Genbank	2995601
Sequence 7714	AC000031	Genbank	4646246
Sequence 7715	AC000036	Genbank	79 <b>5897</b> 9
Sequence 7716	AC000049	Genbank	57 <b>05</b> 995
Sequence 7717	AC000083	Genbank	3810636
Sequence 7718	AC000087	Genbank	7124755
Sequence 7719	AC000088	Genbank	7124834
Sequence 7720	AC000095	Genbank	7940365
Sequence 7721	AC000114	Genbank	1809233
Sequence 7722	AC000134	Genbank	4755212
Sequence 7723	AC000353	Genbank	6970735
Sequence 7724	AC000378	Genbank	2270906
Sequence 7725	AC002054	Genbank	2076723
Sequence 7726	AC002066	Genbank	2076721
Sequence 7727	AC002094	Genbank	2155224
Sequence 7728	AC002105	Genbank	2288973
Sequence 7729	AC002117	Genbank	2281075
Sequence 7730	AC002347	Genbank	2828783
Sequence 7731	AC002377	Genbank	2275192
Sequence 7732	AC002401	Genbank	2599240
Sequence 7733	AC002401	Genbank	6478944
Sequence 7734	AC002522	Genbank	7124014
23443.100 / / 04	, 10002022	Centralik	1 124014

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Sequence 7735	AC002550	Genbank	2570261
Sequence 7736	AC002553	Genbank	3126783
Sequence 7737	AC003010	Genbank	2981254
Sequence 7738	AC003025	Genbank	3337308
Sequence 7739	AC003031	Genbank	2570227
Sequence 7740	AC003036	Genbank	2935595
Sequence 7741	AC003098	Genbank	2822155
Sequence 7742	AC003103	Genbank	2842782
Sequence 7743	AC003663	Genbank	3097871
Sequence 7744	AC003665	Genbank	3808091
Sequence 7745	AC003687	Genbank	2828778
	AC003007	Genbank	4262275
Sequence 7746	AC003963	Genbank	2995604
Sequence 7747		Genbank	2781374
Sequence 7748	AC004016 AC004017	Genbank	2795792
Sequence 7749		Genbank	3219330
Sequence 7750	AC004020		2822164
Sequence 7751	AC004080	Genbank	2822156
Sequence 7752	AC004084	Genbank	
Sequence 7753	AC004098	Genbank	3097872
Sequence 7754	AC004126	Genbank	3242744
Sequence 7755	AC004150	Genbank	4585948
Sequence 7756	AC004228	Genbank	4263838
Sequence 7757	AC004234	Genbank	2914672
Sequence 7758	AC004242	Genbank	3132845
Sequence 7759	AC004253	Genbank	3169206
Sequence 7760	AC004263	Genbank	2935616
Sequence 7761	AC004382	Genbank	3252819
Sequence 7762	AC004410	Genbank	2959558
Sequence 7763	AC004472	Genbank	29 <b>84582</b>
Sequence 7764	AC004475	Genbank	29 <b>88396</b>
Sequence 7765	AC004500	Genbank	2996639
Sequence 7766	AC004522	Genbank	3006227
Sequence 7767	AC004526	Genbank	37 <b>7903</b> 5
Sequence 7768	AC004551	Genbank	3309005
Sequence 7769	AC00-583	Genbank	3293210
Sequence 7770	ACC04696	Genbank	3135391
Sequence 7771	AC004703	Genbank	3228509
Sequence 7772	AC004707	Genbank	3249127
Sequence 7773	AC004757	Genbank	3341707
Sequence 7774	AC004770	Genbank	3212836
Sequence 7775	AC004771	Genbank	3668108
Sequence 7776	AC004801	Genbank	4204244
Sequence 7777	AC004812	Genbank	3970970
Sequence 7778	AC004821	Genbank	4753291
Sequence 7779	AC004823	Genbank	4156207
Sequence 7780	AC004835	Genbank	45081 <b>54</b>
Sequence 7781	AC004854	Genbank	4827328
Sequence 7782	AC004857	Genbank	3694665
Sequence 7783	AC004882	Genbank	5523837
Sequence 7784	AC004893	Genbank	3694662
Sequence 7785	AC0048938	Genbank	3213059
	AC004936 AC004966	Genbank	5306307
Sequence 7786		Genbank	5441941
Sequence 7787	AC004997	Genbank	5649384
Sequence 7788	AC005032	Genbank	4827310
Sequence 7789	AC005037		4508118
Sequence 7790	AC005041	Genbank	4000110

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Sequence 7791	AC005042	Genbank	4156138
Sequence 7792	AC005058	Genbank	4156135
Sequence 7793	AC005071	Genbank	4508112
Sequence 7794	AC005081	Genbank	6560914
Sequence 7795	AC005088	Genbank	4753221
Sequence 7796	AC005104	Genbank	4218027
Sequence 7797	AC005192	Genbank	3264575
Sequence 7798	AC005214	Genbank	3282166
Sequence 7799	AC005215	Genbank	6094661
Sequence 7800	AC005253	Genbank	3288886
Sequence 7801	AC005318	Genbank	3885345
Sequence 7802	AC005316	Genbank	3366582
Sequence 7803	AC005325	Genbank	3366581
Sequence 7804	AC005339		
Sequence 7805		Genbank	3355454
	AC005366	Genbank	3367508
Sequence 7806	AC005369	Genbank	3367505
Sequence 7807	AC005392	Genbank	3399669
Sequence 7808	AC005480	Genbank	4835818
Sequence 7809	AC005488	Genbank	5836194
Sequence 7810	AC005516	Genbank	3900835
Sequence 7811	AC005519	Genbank	5091653
Sequence 7812	AC005523	Genbank	3451032
Sequence 7813	AC005531	Genbank	4153875
Sequence 7814	AC005538	Genbank	4508129
Sequence 7815	AC005546	Genbank	3478635
Sequence 7816	AC005549	Genbank	3598724
Sequence 7817	AC005612	Genbank	3540153
Sequence 7818	AC005629	Genbank	7243877
Sequence 7819	AC005670	Genbank	3688099
Sequence 7820	AC005686	Genbank	3608161
Sequence 7821	AC005726	Genbank	3810672
Sequence 7822	AC005754	Genbank	3688074
Sequence 7823	AC005768	Genbank	6598827
Sequence 7824	AC005789	Genbank	3702281
Sequence 7825	AC005828	Genbank	37 <b>897</b> 13
Sequence 7826	AC005841	Genbank	4731044
Sequence 7827	AC005899	Genbank	3935221
Sequence 7828	AC005921	Genbank	4982556
Sequence 7829	AC005923	Genbank	4309927
Sequence 7830	AC005924	Genbank	4309926
Sequence 7831	AC005988	Genbank	4156128
Sequence 7832	AC006026	Genbank	6042101
Sequence 7833	AC006040	Genbank	4314426
Sequence 7834	AC006042	Genbank	4508120
Sequence 7835	AC006064	Genbank	4572650
Sequence 7836	AC006077	Genbank	3935193
Sequence 7837	AC006088	Genbank	4204701
Sequence 7838	AC006146	Genbank	5836157
Sequence 7839	AC006165	Genbank	3980464
Sequence 7840	AC006241	Genbank	4160141
Sequence 7841	AC006252	Genbank	4309923
Sequence 7842	AC006255	Genbank	5306223
Sequence 7843	AC006299	Genbank	4106997
Sequence 7844	AC006441	Genbank	4760435
Sequence 7845	AC006449	Genbank	6102666
Sequence 7846	AC006478	Genbank	4508145
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Sequence 7847	AC006480	Genbank	6289252
Sequence 7848	AC006487	Genbank	4314418
Sequence 7849	AC006512	Genbank	4926863
S∈quence 7850	AC006987	Genbank	5123989
Sequence 7851	AC007022	Genbank	4753292
Secuence 7852	AC007040	Genbank	4371305
Sequence 7853	AC007172	Genbank	4731066
Secuence 7854	AC007308	Genbank	5903110
Sequence 7855	AC007347	Genbank	6806840
Sequence 7856	AC007381	Genbank	6604532
Sequence 7857	AC007393	Genbank	5306300
Sequence 7858	AC007401	Genbank	6560920
Sequence 7859	AC007617	Genbank	5230397
Sequence 7860	AC007681	Genbank	5836156
Sequence 7861	AC007686	Genbank	6466519
Sequence 7862	AC007688	Genbank	5815499
Sequence 7863	AC007842	Genbank	
Sequence 7864	AC007845	Genbank	5080755 7109495
Sequence 7865	AC008041	Genbank	
Sequence 7866	AC008132	Genbank	5649181 7025879
Sequence 7867	AC008134	Genbank	
Sequence 7868	AC008990	Genbank	5801655
Sequence 7869	AC009330	Genbank	7209394 6553960
Sequence 7870	AC009514	Genbank	6016667
Sequence 7871	AC010077	Genbank	5870275
Sequence 7872	AC011311	Genbank	6850247
Sequence 7873	AC012085	Genbank	6634755
Sequence,7874	AC016026	Genbank	7940357
Sequence 7875	AF000364	Genbank	2697102
Sequence 7876	AF001433	Genbank	2529704
Sequence 7877	AF001435	Genbank	2529708
Sequence 7878	AF001687	Genbank	2653735
Sequence 7879	AF001977	Genbank	2914738
Sequence 7880	AF002020	Genbank	2276462
Sequence 7881	AF002163	Genbank	2290769
Sequence 7882	AF002223	Genbank	4079670
Sequence 7883	AF002697	Genbank	2511528
Sequence 7884	AF002715	Genbank	2352276
Sequence 7885	AF002994	Genbank	2121297
Sequence 7886	AF002996	Genbank	2121299
Sequence 7887	AF004562	Genbank	3041872
Sequence 7888	AF005067	Genbank	6979018
Sequence 7889	AF005258	Genbank	2627428
Sequence 7890	AF005361	Genbank	2343115
Sequence 7891	AF006043	Genbank	2674061
Sequence 7892	AF006083	Genbank	2282031
Sequence 7893	AF006087	Genbank	2282039
Sequence 7894	AF006515	Genbank	2645432
Secuence 7895	AF007170	Genbank	2865 <b>251</b>
Sequence 7896	AF007544	Genbank	2970122
Sequence 7897	AF007791	Genbank	3779196
Sequence 7898	AF007871	Genbank	2358278
Sequence 7899	AF008303	Genbank	2580619
Sequence 7900	AF010187	Genbank	7025910
Sequence 7901	AF012872	Genbank	2326226
Sequence 7902	AF013970	Genbank	2801421

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Sequence 7903	AF014398	Genbank	8148061
Sequence 7904	AF015044	Genbank	4102712
Sequence 7905	AF015913	Genbank	2323409
Sequence 7906	AF016050	Genbank	2978559
Sequence 7907	AF016266	Genbank	252 <b>9562</b>
Sequence 7908	AF017790	Genbank	2501872
Sequence 7909	AF020038	Genbank	3641397
Sequence 7910	AF020736	Genbank	3450954
Sequence 7911	AF020833	Genbank	2460199
Sequence 7912	AF021336	Genbank	2541972
Sequence 7913	AF021351	Genbank	2460207
Sequence 7914	AF022229	Genbank	2809382
Sequence 7915	AF023268	Genbank	2564910
Sequence 7916	AF023462	Genbank	2564670
Sequence 7917	AF023676	Genbank	3211721
Sequence 7918	AF025439	Genbank	2815605
Sequence 7919	AF025999	Genbank	2570 <b>85</b> 3
Sequence 7920	AF026166	Genbank	
Sequence 7921	AF026292	Genbank	4090 <b>928</b>
Secuence 7922	AF026548		2559009
Sequence 7923	AF027299	Genbank Genbank	2583172
Sequence 7924	AF027299 AF027302	Genbank	2739095
Sequence 7925	AF029308		2522533
Sequence 7926	AF031416	Genbank	2564750
Sequence 7927	AF032456	Genbank	3213216
Sequence 7928	AF034373	Genbank	3004908
Sequence 7929	AF034373 AF034756	Genbank	38 <b>20483</b>
Sequence 7930	AF034759	Genbank	2654138
Sequence 7931	AF034759 AF035191	Genbank	2653648
Sequence 7932	AF035191 AF035286	Genbank	4104352
Sequence 7933	AF035295	Genbank	2661038
Sequence 7934	AF035296	Genbank Genbank	266 <b>1050</b>
Sequence 7935	AF036536	Genbank	2661052
Sequence 7936	AF036682	Genbank	4104518
Sequence 7937	AF037448	Genbank	2707821
Sequence 7938	AF037643	Genbank	3037012
Sequence 7939	AF037545	Genbank	2746542
Sequence 7940	AF038318		2795917
Sequence 7941	AF038391	Genbank Genbank	270 <b>8698</b>
Sequence 7942	AF038392	Genbank Genbank	3123905
Sequence 7943	AF038554	Genbank	31 <b>23907</b> 289 <b>5558</b>
Sequence 7944	AF039703	Genbank	
Sequence 7945	AF040707	Genbank	278 <b>9655</b>
Sequence 7946	AF040966	Genbank	368 <b>8794</b>
Sequence 7947	AF041004	Genbank	279 <b>2367</b> 433 <b>5800</b>
Sequence 7948	AF041248	Genbank	
Sequence 7949	AF041259	Genbank	324 <b>3117</b> 33 <b>35396</b>
Sequence 7950	AF041259 AF041381		
Sequence 7951	AF041381 AF041483	Genbank	30 <b>80766</b>
Sequence 7951 Sequence 7952	AF041483 AF042084	Genbank	3493528
Sequence 7953	AF042064 AF042166	Genbank	27 <b>92517</b>
Sequence 7953	AF042379	Genbank	329 <b>8596</b>
Sequence 7955	AF042379 AF043045	Genbank	2801700
Sequence 7955	AF043045 AF043324	Genbank	32 <b>82770</b>
	AF043324 AF044127	Genbank	30 <b>05062</b>
Sequence 7957		Genbank	4105189
Sequence 7958	AF044321	Genbank	317 <b>0263</b>

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Sequence 7959	AF044588	Genbank	2865520
Sequence 7960	AF044773	Genbank	3002950
Sequence 7961	AF044954	Genbank	4164441
Sequence 7962	AF044956	Genbank	5326827
	AF045606	Genbank	8277249
Sequence 7963	AF046025	Genbank	4581063
Sequence 7964			2909853
Sequence 7965	AF047181	Genbank	
Sequence 7966	AF047184	Genbank	2909859
Sequence 7967	AF047431	Genbank	5514630
Sequence 7968	AF047432	Genbank	2911479
Sequence 7969	AF047437	Genbank	3335129
Sequence 7970	AF047442	Genbank	33 <b>35139</b>
Sequence 7971	AF047472	Genbank	2921 <b>872</b>
Sequence 7972	AF047599	Genbank	2906227
Sequence 7973	AF049910	Genbank	3435156
Sequence 7974	AF05C078	Genbank	3818465
Sequence 7975	AF050641	Genbank	5326822
Sequence 7975	AF051782	Genbank	2947237
Sequence 7976		Genbank	6851088
Sequence 7977	AF052059		3360413
Sequence 7978	AF052106	Genbank	3360439
Sequence 7979	AF052130	Genbank	
Sequence 7980	AF052155	Genbank	3360466
Sequence 7981	AF052178	Genbank	3360489
Sequence 7982	AF052179	Genbank	3360490
Sequence 7983	AF052182	Genbank	3360494
Sequence 7984	AF052186	Genbank	3360498
Sequence 7985	AF052504	Genbank	6 <b>650080</b>
Sequence 7986	AF053069	Genbank	3 <b>136151</b>
Sequence 7987	AF053074	Genbank	3283044
Sequence 7988	AF054187	Genbank	4092059
Sequence 7989	AF054284	Genbank	4033734
Sequence 7990	AF054990	Genbank	3005703
Sequence 7991	AF055014	Genbank	3005738
Sequence 7991	AF055017	Genbank	3005743
Sequence 7992	AF055022	Genbank	3005750
Sequence 7993		Genbank	3005759
Sequence 7994	AF055029	Genbank	3694919
Sequence 7995	AF057160		3063679
Sequence 7996	AF058392	Genbank	3300089
Sequence 7997	AF058988	Genbank	
Sequence 7998	AF059321	Genbank	3169 <b>563</b>
Sequence 7999	AF061016	Genbank	3127126
Sequence 8000	AF061243	Genbank	3779243
Sequence 8001	AF061738	Genbank	4335940
Sequence 8002	AF061749	Genbank	3372676
Sequence 8003	AF062536	Genbank	3139076
Sequence 8004	AF063308	Genbank	4106355
Sequence 8005	AF064084	Genbank	3135668
Sequence 8006	AF064603	Genbank	3152659
Sequence 8007	AF065389	Genbank	3152702
Sequence 8008	AF067170	Genbank	4894373
Sequence 8009	AF067513	Genbank	3850305
Sequence 6009	AF067656	Genbank	3901271
Sequence 8010		Genbank	4927369
Sequence 8011	AF067972		4321975
Sequence 8012	AF068235	Genbank	
Sequence 8013	AF068754	Genbank	3283408
Sequence 8014	AF069301	Genbank	3193335

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Sequence 8015	AF069984	Genbank	3242977
Sequence 8016	AF070550	Genbank	. 3387913
Sequence 8017	AF070561	Genbank	3387928
Sequence 8018	AF070563	Genbank	3387931
Sequence 8019	AF070578	Genbank	3387950
Sequence 8020	AF070598	Genbank	3387976
Sequence 8021	AF070603	Genbank	3387983
Sequence 8022	AF070663	Genbank	4454701
Sequence 8023	AF071172	Genbank	5107833
Sequence 8024	AF071202	Genbank	3335172
Sequence 8025	AF071309	Genbank	3426319
Sequence 8026	AF071593	Genbank	3249712
Sequence 8027	AF071738	Genbank	4263875
Sequence 8028	AF071748	Genbank	3916211
Sequence 8029	AF072933	Genbank	5305428
Sequence 8030	AF073344	Genbank	5410229
Sequence 8031	AF074331	Genbank	5052074
Sequence 8032	AF075575	Genbank	3600027
Sequence 8033	AF076273	Genbank	4185718
Sequence 8034	AF076678	Genbank	4884560
Sequence 8035	AF076844	Genbank	4019216
Sequence 8036	AF077036	Genbank	4689119
Sequence 8037	AF077042	Genbank	4689131
Sequence 8038	AF077188	Genbank	5565654
Sequence 8039	AF077203	Genbank	4679019
Sequence 8040	AF077367	Genbank	3820534
Sequence 8041	AF078103	Genbank	4218944
Sequence 8042	AF078107	Genbank	3342907
Sequence 8043	AF078671	Genbank	4877563
Sequence 8044	AF078776	Genbank	3511274
Sequence 8045	AF078845	Genbank	5531804
Sequence 8046	AF078847	Genbank	5531808
Sequence 8047	AF078854	Genbank	5531822
Sequence 8048	AF078863	Genbank	5531840
Sequence 8049	AF078866	Genbank	5531846
Sequence 8050	AF080561	Genbank	3746786
Sequence 8051	AF081484	Genbank	3420928
Sequence 8052	AF083068	Genbank	4808551
Sequence 8053	AF083106	Genbank	7555470
Sequence 8054	AF083190	Genbank	3599414
Sequence 8055	AF083246	Genbank	5106786
Sequence 8056	AF083255	Genbank	3435311
Sequence 8057	AF085355	Genbank	5114044
Sequence 8058	AF085357	Genbank	5114048
Sequence 8059	AF085359	Genbank	5114052
Sequence 8060	AF086234	Genbank	3483579
Sequence 8061	AF086280	Genbank	3483625
Sequence 8062	AF086406	Genbank	3483751
Sequence 8063	AF087438	Genbank	3820591
Sequence 8064	AF087986	Genbank	3523192
Sequence 8065	AF088060	Genbank	3523266
Sequence 8066	AF089841	Genbank	4218954
Sequence 8067	AF090426	Genbank	3982903
Sequence 8068	AF090927	Genbank	6690220
Sequence 8069	AF091035	Genbank	6002584
Sequence 8070	AF091073	Genbank	3859983
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Sequence 8071	AF092136	Genbank	5138921
Sequence 8072	AF092565	Genbank	3661609
Sequence 8073	AF093239	Genbank	3661615
Sequence 8074	AF094516	Genbank	3820613
Sequence 8075	AF095192	Genbank	4322819
Sequence 8076	AF095448	Genbank	4063889
		Genbank	4185790
Sequence 8077	AF095890		4454849
Sequence 8078	AF096160	Genbank	
Sequence 8079	AF097514	Genbank	4808600
Sequence 8080	AF098865	Genbank	4204674
Sequence 8081	AF098948	Genbank	5456746
Sequence 8082	AF100615	Genbank	4808624
Sequence 8083	AF100755	Genbank	5410 <b>295</b>
Sequence 8084	AF100756	Genbank	5410 <b>297</b>
Sequence 8085	AF103800	Genbank	604 <b>8967</b>
Sequence 8086	AF104913	Genbank	3941723
Sequence 8087	AF105366	Genbank	5106522
Sequence 8088	AF106798	Genbank	5805207
Sequence 8089	AF107405	Genbank	5531903
Sequence 8090	AF107406	Genbank	5531905
Sequence 8091	AF110184	Genbank	5565858
Sequence 8092	AF111168	Genbank	4186181
Sequence 8093	AF111853	Genbank	6560625
	AF112227	Genbank	4545218
Sequence 8094	AF113019	Genbank	6642761
Sequence 8095	AF1131129	Genbank	6523820
Sequence 8096		Genbank	5326801
Sequence 8097	AF113132	Genbank	6650673
Sequence 8098	AF113249	Genbank	5852417
Sequence 8099	AF113251	Genbank	6855636
Sequence 8100	AF113702		5006438
Sequence 8101	AF116241	Genbank	5732617
Sequence 8102	AF117888	Genbank	4235636
Sequence 8103	AF118124	Genbank	5931954
Sequence 8104	AF124145	Genbank	
Sequence 8105	AF125102	Genbank	5106999
Sequence 8106	AF125525	Genbank	4689281
Sequence 8107	AF126021	Genbank	6563273
Sequence 8108	AF126246	Genbank	4406523
Sequence 8109	AF126782	Genbank	6318547
Sequence 8110	AF126799	Genbank	4406527
Sequence 8111	AF127138	Genbank	5922724
Sequence 8112	AF127563	Genbank	4455124
Sequence 8113	AF127761	Genbank	4455128
Sequence 8114	AF127966	Genbank	5532271
Sequence 8115	AF128458	Genbank	7140573
Sequence 8116	AF129930	Genbank	4558509
Sequence 8117	AF131742	Genbank	4406554
Sequence 8118	AF131797	Genbank	4406625
Sequence 8119	AF131802	Genbank	4406633
Sequence 8120	AF131820	Genbank	4406655
Sequence 8121	AF131826	Genbank	4406662
Sequence 8122	AF132033	Genbank	5524202
Sequence 8123	AF132856	Genbank	4809025
Sequence 8124	AF132940	Genbank	4680650
Sequence 8125	AF132946	Genbank	4680 <b>662</b>
Sequence 8126	AF132965	Genbank	4680 <b>700</b>

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Sequence 8127	AF134726	Genbank	4529886
Sequence 8128	AF134803	Genbank	4868362
Sequence 8129	AF136271	Genbank	4732145
Sequence 8130	AF136715	Genbank	5019773
Sequence 8131	AF140710	Genbank	6073848
Sequence 8132	AF141347	Genbank	4929133
Sequence 8133	AF141870	Genbank	5006601
Sequence 8134	AF143536	Genbank	4929219
Sequence 8135	AF145732	Genbank	5579330
Sequence 8136	AF147271	Genbank	5823152
Sequence 8137	AF147331	Genbank	4761682
Sequence 8138	AF147410	Genbank	4761761
Sequence 8139	AF147787	Genbank	5231122
Sequence 8140	AF148513	Genbank	4809268
Sequence 8141	AF149045	Genbank	6010479
Sequence 8142	AF150087	Genbank	5107162
Sequence 8143	AF151018	Genbank	7106757
Sequence 8144	AF151020	Genbank	7106761
Sequence 8145	AF151042	Genbank	7106805
Sequence 8146	AF151051		-
Sequence 8147	AF151091	Genbank	7106823
	AF151109 AF151799	Genbank	6166337
Sequence 8148 Sequence 8149	AF151799 AF151805	Genbank	4929550
•	AF151808	Genbank	4929562
Sequence 8150 Sequence 8151	AF151811	Genbank	4929568
		Genbank	4929574
Sequence 8152	AF151826	Genbank	4929604
Sequence 8153 Sequence 8154	AF151841	Genbank	4929634
	AF151846	Genbank	4929644
Sequence 8155	AF151861	Genbank	4929674
Sequence 8156	AF151869	Genbank	4929690
Sequence 8157	AF151886	Genbank	4929724
Sequence 8158	AF151902	Genbank	4929756
Sequence 8159	AF151904	Genbank	4929760
Sequence 8160	AF152961	Genbank	5499740
Sequence 8161	AF155095	Genbank	5360084
Sequence 8162	AF155110	Genbank	5360114
Sequence 8163	AF155238	Genbank	9186822
Sequence 8164	AF156165	Genbank	7145093
Sequence 8165	AF157378	Genbank	5714631
Sequence 8166	AF159295	Genbank	5714635
Sequence 8167	AF161371	Genbank	6841155
Sequence 8168	AF161417	Genbank	6841247
Sequence 8169	AF161448	Genbank	6841309
Sequence 8170	AF161455	Genbank	6841323
Sequence 8171	AF161458	Genbank	6841439
Sequence 8172	AF161485	Genbank	6841493
Sequence 8173	AF161491	Genbank	6841505
Sequence 8174	AF161493	Genbank	6841509
Sequence 8175	AF161511	Genbank	6841545
Sequence 8176	AF161530	Genbank	6841583
Sequence 8177	AF163441	Genbank	6969979
Sequence 8178	AF165967	Genbank	5 <b>802826</b>
Sequence 8179	AF167572	Genbank	6164703
Sequence 8180	AF167706	Genbank	6979310
Sequence 8181	AF168787	Genbank	7239175
Sequence 8182	AF168956	Genbank	5 <b>702387</b>

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Sequence 8183	AF180682	Genbank	6684524
Sequence 8184	AF181265	Genbank	7212810
Sequence 8185	AF182316	Genbank	6731234
Sequence 8186	AF187320	Genbank	6164847
Sequence 8187	AF187554	Genbank	6653225
Sequence 8188	AF190167	Genbank	6456117
	AF190725		
Sequence 8189		Genbank	6118538
Sequence 8190	AF191339	Genbank	6180012
Sequence 8191	AF191340	Genbank	6180014
Sequence 8192	AF192968	Genbank	6449082
Sequence 8193	AF196779	Genbank	6180170
Sequence 8194	AF196969	Genbank	6289073
Sequence 8195	AF198444	Genbank	7025338
Sequence 8196	AF198487	Genbank	6941955
Sequence 8197	AF199596	Genbank	6842049
Sequence 8198	AF203815	Genbank	697 <b>9641</b>
Sequence 8199	AF203978	Genbank	6503203
Sequence 8200	AF207907	Genbank	6653648
Sequence 8201	AF212162	Genbank	6665785
Sequence 8202	AF220417	Genbank	7678700
Sequence 8203	AF223408	Genbank	6911266
Sequence 8204	AF224669	Genbank	7012905
Sequence 8205	AF227218		7159279
,		Genbank Genbank	
Sequence 8206	AJ000186		2463195
Sequence 8207	AJC00332	Genbank	2274967
Sequence 8208	AJ000644	Genbank	2695707
Sequence 8209	AJ001050	Genbank	2832345
Sequence 8210	AJ001443	Genbank	6006514
Sequence 8211	AJC02955	Genbank	2632122
Sequence 8212	AJ004832	Genbank	2982500
Sequence 8213	AJ006291	Genbank	380 <b>5937</b>
Sequence 8214	AJ007398	Genbank	3668140
Sequence 8215	AJ007509	Genbank	3 <b>319955</b>
Sequence 8216	AJ007670	Genbank	3 <b>560123</b>
Sequence 8217	AJ010597	Genbank	3 <b>559873</b>
Sequence 8218	AJ011930	Genbank	3859769
Sequence 8219	AJ012463	Genbank	3850049
Sequence 8220	AJ012499	Genbank	54 <b>41359</b>
Sequence 8221	AJ131186	Genbank	5 <b>689737</b>
Sequence 8222	AJ223366	Genbank	6691913
Sequence 8223	AJ224442	Genbank	2911586
Sequence 8224	AJ245416	Genbank	57 <b>01851</b>
Sequence 8225	AJ250475	Genbank	8919097
Sequence 8226	AJ275986	Genbank	7228110
Sequence 8227	AK000056	Genbank	7019892
Sequence 8228	AK000161	Genbank	7020067
Sequence 8229	AK000181	Genbank	7020098
Sequence 8230	AK000221	Genbank	7020163
Sequence 8231	AK000228	Genbank	7020174
Sequence 8232	AK000226	Genbank	7020174
Sequence 8233	AK000284	Genbank	7020221
	AK000286	Genbank	7020264
Sequence 8234			
Sequence 8235	AK000298	Genbank	7020286
Sequence 8236	AK000301	Genbank	7020291
Sequence 8237	AK000357	Genbank	7020388
Sequence 8238	AK000361	Genbank	7020395

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Sequence 8239         AKC00371         Genbank         7020417           Sequence 8241         AKC000398         Genbank         7020457           Sequence 8242         AK000407         Genbank         7020474           Sequence 8244         AK000436         Genbank         7020522           Sequence 8245         AK000498         Genbank         7020586           Sequence 8246         AK000505         Genbank         7020628           Sequence 8247         AK000543         Genbank         7020709           Sequence 8249         AK000558         Genbank         7020734           Sequence 8250         AK000560         Genbank         7020738           Sequence 8251         AK000560         Genbank         7020738           Sequence 8251         AK000660         Genbank         7020738           Sequence 8253         AK000661         Genbank         7020848           Sequence 8254         AK000661         Genbank         7020882           Sequence 8255         AK000661         Genbank         702099           Sequence 8256         AK000714         Genbank         702099           Sequence 8256         AK000745         Genbank         7021014           Sequence				
Sequence 8241         AKC00038         Genbank         7020459           Sequence 8242         AK000407         Genbank         7020474           Sequence 8244         AK000471         Genbank         7020586           Sequence 8245         AK000498         Genbank         7020586           Sequence 8246         AK000505         Genbank         70207060           Sequence 8247         AK000543         Genbank         7020710           Sequence 8248         AK000554         Genbank         7020734           Sequence 8250         AK000550         Genbank         7020734           Sequence 8251         AK000560         Genbank         7020734           Sequence 8251         AK000660         Genbank         7020734           Sequence 8251         AK000660         Genbank         7020709           Sequence 8253         AK000661         Genbank         7020882           Sequence 8254         AK000661         Genbank         7020909           Sequence 8256         AK000714         Genbank         7020909           Sequence 8256         AK000714         Genbank         7021014           Sequence 8258         AK000745         Genbank         7021025           Sequen	Sequence 8239	AK000371	Genbank	
Sequence 8242         AK000407         Genbank         7020474           Sequence 8243         AK000436         Genbank         7020522           Sequence 8244         AK000471         Genbank         7020586           Sequence 8246         AK000505         Genbank         7020640           Sequence 8248         AK000543         Genbank         7020709           Sequence 8248         AK000558         Genbank         7020734           Sequence 8250         AK000559         Genbank         7020738           Sequence 8251         AK000650         Genbank         7020734           Sequence 8251         AK000650         Genbank         7020796           Sequence 8251         AK000651         Genbank         7020882           Sequence 8254         AK000661         Genbank         7020882           Sequence 8254         AK000664         Genbank         7020909           Sequence 8256         AK000739         Genbank         7021014           Sequence 8257         AK000739         Genbank         7021025           Sequence 8258         AK000740         Genbank         7021025           Sequence 8261         AK000759         Genbank         7021025           Sequenc		AK0 <b>00373</b>	Genbank	7020417
Sequence 8243         AK000436         Genbank         7020522           Sequence 8244         AK000471         Genbank         7020586           Sequence 8245         AK000505         Genbank         7020640           Sequence 8247         AK000543         Genbank         7020709           Sequence 8249         AK000563         Genbank         7020734           Sequence 8250         AK000560         Genbank         7020738           Sequence 8251         AK000560         Genbank         7020738           Sequence 8251         AK000595         Genbank         7020796           Sequence 8253         AK000651         Genbank         7020848           Sequence 8255         AK000661         Genbank         7020882           Sequence 8255         AK000664         Genbank         7020901           Sequence 8255         AK000714         Genbank         7020905           Sequence 8255         AK000739         Genbank         7021014           Sequence 8258         AK000740         Genbank         7021014           Sequence 8259         AK000757         Genbank         7021043           Sequence 8261         AK000759         Genbank         7021043           Sequenc	Sequence 8241	AK00 <b>039</b> 8	Genbank	7020459
Sequence 8243         AK000436         Genbank         7020522           Sequence 8244         AK000474         Genbank         7020586           Sequence 8245         AK000505         Genbank         7020640           Sequence 8247         AK000543         Genbank         7020710           Sequence 8249         AK000543         Genbank         7020710           Sequence 8249         AK000560         Genbank         7020734           Sequence 8251         AK000560         Genbank         7020738           Sequence 8251         AK000595         Genbank         7020796           Sequence 8253         AK000651         Genbank         7020848           Sequence 8253         AK000651         Genbank         7020848           Sequence 8255         AK000664         Genbank         7020901           Sequence 8255         AK000664         Genbank         7020901           Sequence 8255         AK000714         Genbank         7021014           Sequence 8258         AK000740         Genbank         7021014           Sequence 8259         AK000757         Genbank         7021043           Sequence 8261         AK000759         Genbank         7021043           Sequenc	Sequence 8242	AK00 <b>0407</b>	Genbank	7020474
Sequence 8244         AK000471         Genbank         7020586           Sequence 8245         AK000498         Genbank         7020628           Sequence 8246         AK000505         Genbank         7020640           Sequence 8247         AK000543         Genbank         7020710           Sequence 8248         AK000560         Genbank         7020734           Sequence 8250         AK000560         Genbank         7020736           Sequence 8251         AK000695         Genbank         7020796           Sequence 8252         AK000630         Genbank         7020848           Sequence 8253         AK000661         Genbank         7020848           Sequence 8255         AK000664         Genbank         7020901           Sequence 8255         AK000669         Genbank         7020909           Sequence 8256         AK000714         Genbank         7020975           Sequence 8257         AK000739         Genbank         7021016           Sequence 8258         AK000740         Genbank         7021025           Sequence 8261         AK000759         Genbank         7021046           Sequence 8263         AK000759         Genbank         7021103           Sequenc	Sequence 8243	AK000436	Genbank	
Sequence 8245         AKC00498         Genbank         7020628           Sequence 8246         AKC000543         Genbank         7020640           Sequence 8247         AK000543         Genbank         7020710           Sequence 8249         AK000558         Genbank         7020734           Sequence 8250         AK000560         Genbank         7020734           Sequence 8251         AK000595         Genbank         7020796           Sequence 8252         AK000630         Genbank         7020848           Sequence 8253         AK000661         Genbank         7020901           Sequence 8254         AK000669         Genbank         7020903           Sequence 8255         AK000669         Genbank         7020903           Sequence 8256         AK000714         Genbank         7021014           Sequence 8255         AK000739         Genbank         7021015           Sequence 8255         AK000740         Genbank         7021015           Sequence 8261         AK000757         Genbank         7021043           Sequence 8261         AK000759         Genbank         7021045           Sequence 8263         AK000846         Genbank         7021183           Sequen		AK000471	Genbank	7020586
Sequence 8246         AKC00505         Genbank         7020640           Sequence 8247         AKC000543         Genbank         7020709           Sequence 8248         AKC000544         Genbank         7020710           Sequence 8249         AKC000560         Genbank         7020738           Sequence 8250         AKC000595         Genbank         7020798           Sequence 8251         AKC000630         Genbank         7020848           Sequence 8253         AKC000651         Genbank         7020848           Sequence 8254         AKC000664         Genbank         7020901           Sequence 8255         AKC000669         Genbank         7020909           Sequence 8256         AKC000749         Genbank         7020909           Sequence 8257         AKC000739         Genbank         7021014           Sequence 8258         AKC000740         Genbank         7021014           Sequence 8259         AKC000757         Genbank         7021045           Sequence 8261         AKC000759         Genbank         7021046           Sequence 8263         AKC000849         Genbank         7021103           Sequence 8265         AKC000946         Genbank         7021980				
Sequence 8247         AK000543         Genbank         7020709           Sequence 8248         AK000544         Genbank         7020710           Sequence 8259         AK000560         Genbank         7020738           Sequence 8251         AK000595         Genbank         7020796           Sequence 8252         AK000630         Genbank         7020848           Sequence 8253         AK000661         Genbank         7020901           Sequence 8255         AK000669         Genbank         7020909           Sequence 8256         AK000714         Genbank         7020909           Sequence 8256         AK000714         Genbank         7020909           Sequence 8256         AK000740         Genbank         7021016           Sequence 8257         AK000740         Genbank         7021016           Sequence 8261         AK000757         Genbank         7021043           Sequence 8261         AK000759         Genbank         7021046           Sequence 8263         AK000959         Genbank         7021043           Sequence 8264         AK000946         Genbank         7021183           Sequence 8265         AK000980         Genbank         7021928           Sequenc		AK000505		
Sequence 8248         AK000554         Genbank         7020710           Sequence 8250         AK000558         Genbank         7020734           Sequence 8251         AK000595         Genbank         7020796           Sequence 8251         AK000630         Genbank         7020848           Sequence 8253         AK000651         Genbank         7020848           Sequence 8254         AK000669         Genbank         7020909           Sequence 8256         AK000714         Genbank         7020909           Sequence 8256         AK000740         Genbank         7020909           Sequence 8256         AK000749         Genbank         7021014           Sequence 8258         AK000740         Genbank         7021014           Sequence 8258         AK000745         Genbank         7021025           Sequence 8261         AK000757         Genbank         7021043           Sequence 8261         AK000759         Genbank         7021148           Sequence 8263         AK000859         Genbank         7021183           Sequence 8264         AK000982         Genbank         7021980           Sequence 8265         AK000998         Genbank         7021980           Sequenc				
Sequence 8249         AK000558         Genbank         7020734           Sequence 8251         AK000560         Genbank         7020738           Sequence 8251         AK000630         Genbank         7020848           Sequence 8253         AK000651         Genbank         7020882           Sequence 8254         AK000664         Genbank         7020901           Sequence 8255         AK000669         Genbank         7020907           Sequence 8256         AK000714         Genbank         7021014           Sequence 8257         AK000739         Genbank         7021014           Sequence 8258         AK000740         Genbank         7021015           Sequence 8260         AK000757         Genbank         7021025           Sequence 8261         AK000759         Genbank         7021046           Sequence 8263         AK000859         Genbank         7021103           Sequence 8264         AK000946         Genbank         7021193           Sequence 8265         AK000982         Genbank         7021928           Sequence 8266         AK001190         Genbank         7021980           Sequence 8268         AK001120         Genbank         7022198           Sequenc				
Sequence 8250         AK000560         Genbank         7020738           Sequence 8251         AK000595         Genbank         7020796           Sequence 8252         AK000630         Genbank         7020848           Sequence 8254         AK000661         Genbank         7020901           Sequence 8255         AK000669         Genbank         7020909           Sequence 8256         AK000714         Genbank         7020975           Sequence 8257         AK000739         Genbank         7021016           Sequence 8258         AK000740         Genbank         7021016           Sequence 8260         AK000757         Genbank         7021043           Sequence 8261         AK000759         Genbank         7021046           Sequence 8262         AK000759         Genbank         7021043           Sequence 8263         AK000859         Genbank         7021103           Sequence 8264         AK000982         Genbank         7021183           Sequence 8265         AK000982         Genbank         7021928           Sequence 8266         AK000990         Genbank         7021928           Sequence 8268         AK001112         Genbank         7022193           Sequenc				
Sequence 8251         AK000595         Genbank         7020796           Sequence 8253         AK000630         Genbank         7020848           Sequence 8253         AK000661         Genbank         7020882           Sequence 8255         AK000664         Genbank         7020909           Sequence 8256         AK000714         Genbank         7020975           Sequence 8257         AK000739         Genbank         7021014           Sequence 8258         AK000740         Genbank         7021016           Sequence 8260         AK000757         Genbank         7021025           Sequence 8261         AK000759         Genbank         7021043           Sequence 8263         AK000759         Genbank         7021045           Sequence 8263         AK000859         Genbank         7021103           Sequence 8264         AK000946         Genbank         7021928           Sequence 8265         AK000990         Genbank         7021928           Sequence 8266         AK000199         Genbank         7021928           Sequence 8269         AK001112         Genbank         7022139           Sequence 8269         AK001123         Genbank         7022170           Sequenc				
Sequence 8252         AK000630         Genbank         7020884           Sequence 8253         AK000661         Genbank         7020802           Sequence 8255         AK000669         Genbank         7020901           Sequence 8256         AK000714         Genbank         70209075           Sequence 8257         AK000739         Genbank         7021014           Sequence 8258         AK000740         Genbank         7021015           Sequence 8260         AK000757         Genbank         7021043           Sequence 8261         AK000759         Genbank         7021043           Sequence 8263         AK000759         Genbank         7021043           Sequence 8263         AK000859         Genbank         7021043           Sequence 8263         AK000859         Genbank         7021928           Sequence 8264         AK000946         Genbank         7021980           Sequence 8265         AK000990         Genbank         7021980           Sequence 8266         AK0001990         Genbank         7022198           Sequence 8267         AK001120         Genbank         7022198           Sequence 8271         AK001184         Genbank         7022170           Seque				
Sequence 8253         AK000651         Genbank         7020882           Sequence 8254         AK000664         Genbank         7020909           Sequence 8255         AK000669         Genbank         7020909           Sequence 8256         AK000714         Genbank         7021014           Sequence 8258         AK000740         Genbank         7021016           Sequence 8259         AK000745         Genbank         7021043           Sequence 8261         AK000759         Genbank         7021043           Sequence 8261         AK000759         Genbank         7021043           Sequence 8263         AK000859         Genbank         7021043           Sequence 8263         AK000859         Genbank         7021103           Sequence 8264         AK000946         Genbank         7021928           Sequence 8265         AK000990         Genbank         7021980           Sequence 8266         AK001090         Genbank         702193           Sequence 8266         AK001123         Genbank         7022139           Sequence 8273         AK001184         Genbank         7022213           Sequence 8274         AK001188         Genbank         7022246           Sequence				
Sequence 8254         AK000664         Genbank         7020901           Sequence 8255         AK000669         Genbank         7020909           Sequence 8256         AK000714         Genbank         7020975           Sequence 8257         AK000739         Genbank         7021014           Sequence 8258         AK000740         Genbank         7021015           Sequence 8260         AK000757         Genbank         7021043           Sequence 8261         AK000759         Genbank         7021046           Sequence 8262         AK000759         Genbank         7021103           Sequence 8263         AK000859         Genbank         7021183           Sequence 8263         AK000990         Genbank         7021928           Sequence 8264         AK000990         Genbank         7021992           Sequence 8268         AK001123         Genbank         7022170           Sequence 8268         AK001123         Genbank         7022170           Sequence 8269         AK001123         Genbank         7022170           Sequence 8271         AK001188         Genbank         70222186           Sequence 8273         AK001262         Genbank         70222479           Seque				
Sequence 8255         AK000669         Genbank         7020909           Sequence 8256         AK000714         Genbank         7020975           Sequence 8257         AK000740         Genbank         7021016           Sequence 8258         AK000745         Genbank         7021025           Sequence 8260         AK000757         Genbank         7021043           Sequence 8261         AK000759         Genbank         7021046           Sequence 8263         AK000859         Genbank         7021103           Sequence 8264         AK000989         Genbank         7021183           Sequence 8265         AK000982         Genbank         7021928           Sequence 8264         AK000990         Genbank         7021980           Sequence 8265         AK001090         Genbank         7021980           Sequence 8266         AK001090         Genbank         7021980           Sequence 8267         AK001190         Genbank         7022139           Sequence 8268         AK001112         Genbank         7022170           Sequence 8273         AK001184         Genbank         7022278           Sequence 8271         AK001262         Genbank         7022246           Sequenc				
Sequence 8256         AK000714         Genbank         7020975           Sequence 8257         AK000739         Genbank         7021014           Sequence 8258         AK000740         Genbank         7021016           Sequence 8259         AK000745         Genbank         7021025           Sequence 8261         AK000759         Genbank         7021043           Sequence 8261         AK000799         Genbank         7021103           Sequence 8263         AK000859         Genbank         7021193           Sequence 8264         AK000946         Genbank         7021928           Sequence 8265         AK000990         Genbank         7021992           Sequence 8266         AK000990         Genbank         7021992           Sequence 8267         AK001090         Genbank         7022170           Sequence 8269         AK001123         Genbank         7022170           Sequence 8270         AK001184         Genbank         7022279           Sequence 8271         AK001188         Genbank         7022279           Sequence 8273         AK001262         Genbank         7022240           Sequence 8273         AK001264         Genbank         702240           Sequence				
Sequence 8257         AK000739         Genbank         7021014           Sequence 8258         AK000740         Genbank         7021016           Sequence 8259         AK000745         Genbank         7021025           Sequence 8260         AK000757         Genbank         7021043           Sequence 8261         AK000759         Genbank         7021043           Sequence 8262         AK000799         Genbank         7021183           Sequence 8263         AK000859         Genbank         7021183           Sequence 8264         AK000982         Genbank         7021928           Sequence 8265         AK000990         Genbank         7021992           Sequence 8267         AK001190         Genbank         7022139           Sequence 8268         AK001112         Genbank         7022170           Sequence 8269         AK001123         Genbank         7022170           Sequence 8271         AK001184         Genbank         7022279           Sequence 8271         AK001188         Genbank         7022240           Sequence 8273         AK001262         Genbank         702240           Sequence 8274         AK001281         Genbank         7022419           Sequence				
Sequence 8258         AK000740         Genbank         7021016           Sequence 8259         AK000745         Genbank         7021025           Sequence 8260         AK000757         Genbank         7021043           Sequence 8261         AK000759         Genbank         7021046           Sequence 8262         AK000799         Genbank         7021103           Sequence 8263         AK0008859         Genbank         7021198           Sequence 8264         AK000982         Genbank         7021928           Sequence 8265         AK000990         Genbank         7021992           Sequence 8266         AK001090         Genbank         7022179           Sequence 8268         AK001112         Genbank         7022170           Sequence 8269         AK001123         Genbank         7022170           Sequence 8270         AK001184         Genbank         7022279           Sequence 8271         AK001188         Genbank         7022240           Sequence 8271         AK001262         Genbank         7022409           Sequence 8274         AK001264         Genbank         7022419           Sequence 8276         AK001277         Genbank         7022419           Sequen				
Sequence 8259         AK000745         Genbank         7021025           Sequence 8260         AK000757         Genbank         7021043           Sequence 8261         AK000759         Genbank         7021046           Sequence 8262         AK000799         Genbank         7021103           Sequence 8263         AK000859         Genbank         7021183           Sequence 8264         AK000982         Genbank         7021928           Sequence 8265         AK000990         Genbank         7021992           Sequence 8266         AK001090         Genbank         7022139           Sequence 8268         AK001123         Genbank         7022170           Sequence 8269         AK001123         Genbank         70222170           Sequence 8270         AK001184         Genbank         70222170           Sequence 8271         AK001188         Genbank         70222279           Sequence 8271         AK001188         Genbank         70222406           Sequence 8273         AK001262         Genbank         7022406           Sequence 8274         AK001269         Genbank         7022409           Sequence 8275         AK001277         Genbank         7022437           Seq				
Sequence 8260         AK000757         Genbank         7021043           Sequence 8261         AK000759         Genbank         7021046           Sequence 8262         AK000799         Genbank         7021103           Sequence 8263         AK000859         Genbank         7021183           Sequence 8264         AK000946         Genbank         7021980           Sequence 8265         AK000990         Genbank         7021992           Sequence 8266         AK001090         Genbank         7022139           Sequence 8268         AK001112         Genbank         7022170           Sequence 8269         AK0011123         Genbank         7022170           Sequence 8270         AK001184         Genbank         7022279           Sequence 8271         AK001188         Genbank         7022279           Sequence 8271         AK001262         Genbank         7022406           Sequence 8273         AK001264         Genbank         7022409           Sequence 8274         AK001269         Genbank         7022409           Sequence 8275         AK001277         Genbank         7022431           Sequence 8276         AK001281         Genbank         7022448           Sequen	•			
Sequence 8261         AK000759         Genbank         7021046           Sequence 8262         AK000799         Genbank         7021103           Sequence 8263         AK000859         Genbank         7021183           Sequence 8264         AK000946         Genbank         7021928           Sequence 8265         AK000990         Genbank         7021992           Sequence 8266         AK000990         Genbank         7022139           Sequence 8268         AK001112         Genbank         7022170           Sequence 8269         AK001112         Genbank         7022170           Sequence 8270         AK001184         Genbank         7022279           Sequence 8271         AK001188         Genbank         7022240           Sequence 8272         AK001262         Genbank         7022406           Sequence 8273         AK001264         Genbank         7022409           Sequence 8274         AK001269         Genbank         7022419           Sequence 8275         AK001277         Genbank         7022431           Sequence 8276         AK001281         Genbank         7022431           Sequence 8278         AK001292         Genbank         7022431           Sequenc				
Sequence 8262         AK000799         Genbank         7021103           Sequence 8263         AK000859         Genbank         7021183           Sequence 8264         AK000946         Genbank         7021928           Sequence 8265         AK000990         Genbank         7021992           Sequence 8266         AK001090         Genbank         7022139           Sequence 8267         AK001190         Genbank         7022170           Sequence 8268         AK001112         Genbank         7022170           Sequence 8269         AK001123         Genbank         7022176           Sequence 8270         AK001184         Genbank         7022279           Sequence 8271         AK001188         Genbank         70222406           Sequence 8272         AK001262         Genbank         7022406           Sequence 8273         AK001264         Genbank         7022406           Sequence 8273         AK001269         Genbank         7022409           Sequence 8274         AK001269         Genbank         7022419           Sequence 8275         AK001277         Genbank         7022437           Sequence 8278         AK001288         Genbank         7022455           Sequen				
Sequence         8263         AK000859         Genbank         7021183           Sequence         8264         AK000946         Genbank         7021928           Sequence         8265         AK000990         Genbank         7021980           Sequence         8266         AK001090         Genbank         7021992           Sequence         8267         AK001090         Genbank         7022170           Sequence         8268         AK001112         Genbank         7022170           Sequence         8269         AK001123         Genbank         7022170           Sequence         8270         AK001184         Genbank         7022276           Sequence         8271         AK001188         Genbank         7022276           Sequence         8272         AK001262         Genbank         7022409           Sequence         8273         AK001269         Genbank         7022409           Sequence         8274         AK001269         Genbank         7022409           Sequence         8275         AK001277         Genbank         7022431           Sequence         8277         AK001281         Genbank         7022437           Sequence				
Sequence 8264         AK000946         Genbank         7021928           Sequence 8265         AK000982         Genbank         7021980           Sequence 8266         AK000990         Genbank         7021992           Sequence 8267         AK001090         Genbank         7022139           Sequence 8268         AK001112         Genbank         7022170           Sequence 8269         AK001123         Genbank         7022170           Sequence 8270         AK001184         Genbank         7022279           Sequence 8271         AK001188         Genbank         7022279           Sequence 8272         AK001262         Genbank         7022406           Sequence 8273         AK001264         Genbank         7022409           Sequence 8274         AK001269         Genbank         7022419           Sequence 8275         AK001277         Genbank         7022431           Sequence 8276         AK001281         Genbank         7022437           Sequence 8277         AK001288         Genbank         7022448           Sequence 8278         AK001292         Genbank         7022448           Sequence 8280         AK001313         Genbank         702250           Sequence				
Sequence 8265         AK000982         Genbank         7021980           Sequence 8266         AK000990         Genbank         7021992           Sequence 8267         AK001090         Genbank         7022139           Sequence 8268         AK001112         Genbank         7022170           Sequence 8269         AK001123         Genbank         7022176           Sequence 8270         AK001184         Genbank         7022279           Sequence 8271         AK001188         Genbank         70222406           Sequence 8272         AK001262         Genbank         7022406           Sequence 8273         AK001264         Genbank         7022409           Sequence 8274         AK001269         Genbank         7022409           Sequence 8275         AK001277         Genbank         7022419           Sequence 8276         AK001281         Genbank         7022431           Sequence 8276         AK001281         Genbank         7022437           Sequence 8277         AK001288         Genbank         7022448           Sequence 8279         AK001313         Genbank         7022455           Sequence 8280         AK001331         Genbank         7022543           Sequen				
Sequence         8266         AK000990         Genbank         7021992           Sequence         8267         AK001090         Genbank         7022139           Sequence         8268         AK001112         Genbank         7022170           Sequence         8269         AK001123         Genbank         7022186           Sequence         8270         AK001184         Genbank         7022279           Sequence         8271         AK001188         Genbank         7022406           Sequence         8272         AK001262         Genbank         7022406           Sequence         8273         AK001264         Genbank         7022409           Sequence         8274         AK001269         Genbank         7022419           Sequence         8275         AK001277         Genbank         7022431           Sequence         8276         AK001281         Genbank         7022437           Sequence         8277         AK001288         Genbank         7022448           Sequence         8278         AK001292         Genbank         7022455           Sequence         8281         AK001331         Genbank         7022543           Sequence				
Sequence 8267         AK001090         Genbank         7022139           Sequence 8268         AK001112         Genbank         7022170           Sequence 8269         AK001123         Genbank         7022186           Sequence 8270         AK001184         Genbank         7022279           Sequence 8271         AK001188         Genbank         7022286           Sequence 8272         AK001262         Genbank         7022406           Sequence 8273         AK001264         Genbank         7022409           Sequence 8274         AK001269         Genbank         7022419           Sequence 8275         AK001277         Genbank         7022431           Sequence 8276         AK001281         Genbank         7022437           Sequence 8277         AK001288         Genbank         7022448           Sequence 8278         AK001292         Genbank         7022445           Sequence 8279         AK001313         Genbank         7022455           Sequence 8280         AK001330         Genbank         7022543           Sequence 8281         AK001344         Genbank         7022543           Sequence 8282         AK001363         Genbank         7022575           Sequenc				
Sequence         8268         AK001112         Genbank         7022170           Sequence         8269         AK001123         Genbank         7022186           Sequence         8270         AK001184         Genbank         7022279           Sequence         8271         AK001188         Genbank         7022286           Sequence         8272         AK001262         Genbank         7022406           Sequence         8273         AK001264         Genbank         7022409           Sequence         8274         AK001269         Genbank         7022419           Sequence         8275         AK001277         Genbank         7022431           Sequence         8276         AK001281         Genbank         7022437           Sequence         8277         AK001288         Genbank         7022448           Sequence         8278         AK001292         Genbank         7022448           Sequence         8279         AK001313         Genbank         7022490           Sequence         8281         AK001330         Genbank         7022543           Sequence         8281         AK001363         Genbank         7022575           Sequence				
Sequence 8269         AK001123         Genbank         7022186           Sequence 8270         AK001184         Genbank         7022279           Sequence 8271         AK001188         Genbank         7022286           Sequence 8272         AK001262         Genbank         7022406           Sequence 8273         AK001264         Genbank         7022409           Sequence 8274         AK001269         Genbank         7022419           Sequence 8275         AK001277         Genbank         7022431           Sequence 8276         AK001281         Genbank         7022437           Sequence 8277         AK001288         Genbank         7022448           Sequence 8278         AK001292         Genbank         7022455           Sequence 8280         AK001313         Genbank         7022490           Sequence 8280         AK001330         Genbank         7022543           Sequence 8281         AK001344         Genbank         7022543           Sequence 8282         AK001363         Genbank         7022573           Sequence 8283         AK001364         Genbank         7022575           Sequence 8284         AK001387         Genbank         7022579           Sequenc				
Sequence 8270         AK001184         Genbank         7022279           Sequence 8271         AK001188         Genbank         7022286           Sequence 8272         AK001262         Genbank         7022406           Sequence 8273         AK001264         Genbank         7022409           Sequence 8274         AK001269         Genbank         7022419           Sequence 8275         AK001277         Genbank         7022431           Sequence 8276         AK001281         Genbank         7022437           Sequence 8277         AK001288         Genbank         7022448           Sequence 8278         AK001292         Genbank         7022455           Sequence 8279         AK001313         Genbank         7022490           Sequence 8280         AK001330         Genbank         7022523           Sequence 8281         AK001344         Genbank         7022525           Sequence 8282         AK001363         Genbank         7022575           Sequence 8283         AK001364         Genbank         7022579           Sequence 8284         AK001387         Genbank         7022579           Sequence 8285         AK001392         Genbank         7022625           Sequenc				
Sequence 8271         AK001188         Genbank         7022286           Sequence 8272         AK001262         Genbank         7022406           Sequence 8273         AK001264         Genbank         7022409           Sequence 8274         AK001269         Genbank         7022419           Sequence 8275         AK001277         Genbank         7022431           Sequence 8276         AK001281         Genbank         7022437           Sequence 8277         AK001288         Genbank         7022448           Sequence 8278         AK001292         Genbank         7022455           Sequence 8279         AK001313         Genbank         7022455           Sequence 8280         AK001330         Genbank         7022520           Sequence 8281         AK001344         Genbank         7022523           Sequence 8282         AK001363         Genbank         7022573           Sequence 8283         AK001364         Genbank         7022577           Sequence 8284         AK001366         Genbank         7022579           Sequence 8285         AK001387         Genbank         7022579           Sequence 8286         AK001392         Genbank         7022621           Sequenc			-	
Sequence 8272         AK001262         Genbank         7022406           Sequence 8273         AK001264         Genbank         7022409           Sequence 8274         AK001269         Genbank         7022419           Sequence 8275         AK001277         Genbank         7022431           Sequence 8276         AK001281         Genbank         7022437           Sequence 8277         AK001288         Genbank         7022448           Sequence 8278         AK001292         Genbank         7022455           Sequence 8279         AK001313         Genbank         7022490           Sequence 8280         AK001330         Genbank         7022520           Sequence 8281         AK001344         Genbank         7022543           Sequence 8282         AK001363         Genbank         7022573           Sequence 8283         AK001364         Genbank         7022575           Sequence 8284         AK001366         Genbank         7022575           Sequence 8285         AK001387         Genbank         7022579           Sequence 8286         AK001387         Genbank         7022615           Sequence 8287         AK001395         Genbank         7022625           Sequenc	•			
Sequence 8273         AK001264         Genbank         7022409           Sequence 8274         AK001269         Genbank         7022419           Sequence 8275         AK001277         Genbank         7022431           Sequence 8276         AK001281         Genbank         7022437           Sequence 8277         AK001288         Genbank         7022448           Sequence 8278         AK001292         Genbank         7022455           Sequence 8279         AK001313         Genbank         7022490           Sequence 8280         AK001330         Genbank         7022520           Sequence 8281         AK001344         Genbank         7022573           Sequence 8282         AK001363         Genbank         7022575           Sequence 8283         AK001364         Genbank         7022577           Sequence 8284         AK001364         Genbank         7022577           Sequence 8284         AK001366         Genbank         7022579           Sequence 8285         AK001387         Genbank         7022579           Sequence 8286         AK001392         Genbank         7022621           Sequence 8287         AK001395         Genbank         7022625           Sequenc			_ +	
Sequence 8274         AK001269         Genbank         7022419           Sequence 8275         AK001277         Genbank         7022431           Sequence 8276         AK001281         Genbank         7022437           Sequence 8277         AK001288         Genbank         7022448           Sequence 8278         AK001292         Genbank         7022455           Sequence 8279         AK001313         Genbank         7022490           Sequence 8280         AK001330         Genbank         7022520           Sequence 8281         AK001344         Genbank         7022543           Sequence 8282         AK001363         Genbank         7022573           Sequence 8283         AK001364         Genbank         7022577           Sequence 8284         AK001366         Genbank         7022577           Sequence 8285         AK001387         Genbank         7022579           Sequence 8286         AK001387         Genbank         7022615           Sequence 8287         AK001395         Genbank         7022625           Sequence 8288         AK001410         Genbank         7022625           Sequence 8289         AK001478         Genbank         7022692           Sequenc				
Sequence 8275         AK001277         Genbank         7022431           Sequence 8276         AK001281         Genbank         7022437           Sequence 8277         AK001288         Genbank         7022448           Sequence 8278         AK001292         Genbank         7022455           Sequence 8279         AK001313         Genbank         7022490           Sequence 8280         AK001330         Genbank         7022520           Sequence 8281         AK001344         Genbank         7022543           Sequence 8282         AK001363         Genbank         7022575           Sequence 8283         AK001364         Genbank         7022575           Sequence 8284         AK001366         Genbank         7022577           Sequence 8285         AK001387         Genbank         7022575           Sequence 8286         AK001387         Genbank         7022615           Sequence 8287         AK001392         Genbank         7022621           Sequence 8288         AK001410         Genbank         7022625           Sequence 8289         AK001410         Genbank         7022692           Sequence 8290         AK001478         Genbank         7022760           Sequenc				
Sequence 8276         AK001281         Genbank         7022437           Sequence 8277         AK001288         Genbank         7022448           Sequence 8278         AK001292         Genbank         7022455           Sequence 8279         AK001313         Genbank         7022490           Sequence 8280         AK001330         Genbank         7022520           Sequence 8281         AK001344         Genbank         7022543           Sequence 8282         AK001363         Genbank         7022575           Sequence 8283         AK001364         Genbank         7022577           Sequence 8284         AK001366         Genbank         7022579           Sequence 8285         AK001387         Genbank         7022579           Sequence 8286         AK001392         Genbank         7022621           Sequence 8287         AK001395         Genbank         7022625           Sequence 8288         AK001410         Genbank         7022692           Sequence 8299         AK001478         Genbank         7022760           Sequence 8291         AK001492         Genbank         7022781           Sequence 8292         AK001508         Genbank         7022805           Sequenc				
Sequence 8277         AK001288         Genbank         7022448           Sequence 8278         AK001292         Genbank         7022455           Sequence 8279         AK001313         Genbank         7022490           Sequence 8280         AK001330         Genbank         7022520           Sequence 8281         AK001344         Genbank         7022543           Sequence 8282         AK001363         Genbank         7022575           Sequence 8283         AK001364         Genbank         7022577           Sequence 8284         AK001366         Genbank         7022579           Sequence 8285         AK001387         Genbank         7022579           Sequence 8286         AK001392         Genbank         7022615           Sequence 8287         AK001392         Genbank         7022621           Sequence 8288         AK001395         Genbank         7022625           Sequence 8289         AK001410         Genbank         7022692           Sequence 8290         AK001478         Genbank         7022760           Sequence 8291         AK001492         Genbank         7022781           Sequence 8292         AK001508         Genbank         7022805           Sequenc				
Sequence 8278         AK001292         Genbank         7022455           Sequence 8279         AK001313         Genbank         7022490           Sequence 8280         AK001330         Genbank         7022520           Sequence 8281         AK001344         Genbank         7022543           Sequence 8282         AK001363         Genbank         7022575           Sequence 8283         AK001364         Genbank         7022577           Sequence 8284         AK001366         Genbank         7022579           Sequence 8285         AK001387         Genbank         7022615           Sequence 8286         AK001392         Genbank         7022621           Sequence 8287         AK001395         Genbank         7022625           Sequence 8288         AK001410         Genbank         7022649           Sequence 8289         AK001436         Genbank         7022692           Sequence 8291         AK001478         Genbank         7022760           Sequence 8292         AK001508         Genbank         7022805           Sequence 8293         AK001521         Genbank         7022828				
Sequence 8279         AK001313         Genbank         7022490           Sequence 8280         AK001330         Genbank         7022520           Sequence 8281         AK001344         Genbank         7022543           Sequence 8282         AK001363         Genbank         7022575           Sequence 8283         AK001364         Genbank         7022577           Sequence 8284         AK001366         Genbank         7022579           Sequence 8285         AK001387         Genbank         7022615           Sequence 8286         AK001392         Genbank         7022621           Sequence 8287         AK001395         Genbank         7022625           Sequence 8288         AK001410         Genbank         7022649           Sequence 8289         AK001436         Genbank         7022692           Sequence 8290         AK001478         Genbank         7022760           Sequence 8291         AK001492         Genbank         7022781           Sequence 8292         AK001508         Genbank         7022805           Sequence 8293         AK001521         Genbank         7022828	•			
Sequence 8280         AK001330         Genbank         7022520           Sequence 8281         AK001344         Genbank         7022543           Sequence 8282         AK001363         Genbank         7022575           Sequence 8283         AK001364         Genbank         7022577           Sequence 8284         AK001366         Genbank         7022579           Sequence 8285         AK001387         Genbank         7022615           Sequence 8286         AK001392         Genbank         7022621           Sequence 8287         AK001395         Genbank         7022625           Sequence 8288         AK001410         Genbank         7022649           Sequence 8289         AK001436         Genbank         7022692           Sequence 8290         AK001478         Genbank         7022760           Sequence 8291         AK001492         Genbank         7022781           Sequence 8292         AK001508         Genbank         7022805           Sequence 8293         AK001521         Genbank         7022828				
Sequence 8281         AK001344         Genbank         7022543           Sequence 8282         AK001363         Genbank         7022575           Sequence 8283         AK001364         Genbank         7022577           Sequence 8284         AK001366         Genbank         7022579           Sequence 8285         AK001387         Genbank         7022615           Sequence 8286         AK001392         Genbank         7022621           Sequence 8287         AK001395         Genbank         7022625           Sequence 8288         AK001410         Genbank         7022649           Sequence 8289         AK001436         Genbank         7022692           Sequence 8290         AK001478         Genbank         7022760           Sequence 8291         AK001492         Genbank         7022781           Sequence 8292         AK001508         Genbank         7022805           Sequence 8293         AK001521         Genbank         7022828				
Sequence 8282         AK001363         Genbank         7022575           Sequence 8283         AK001364         Genbank         7022577           Sequence 8284         AK001366         Genbank         7022579           Sequence 8285         AK001387         Genbank         7022615           Sequence 8286         AK001392         Genbank         7022621           Sequence 8287         AK001395         Genbank         7022625           Sequence 8288         AK001410         Genbank         7022649           Sequence 8289         AK001436         Genbank         7022692           Sequence 8290         AK001478         Genbank         7022760           Sequence 8291         AK001492         Genbank         7022781           Sequence 8292         AK001508         Genbank         7022805           Sequence 8293         AK001521         Genbank         7022828				
Sequence 8283         AK001364         Genbank         7022577           Sequence 8284         AK001366         Genbank         7022579           Sequence 8285         AK001387         Genbank         7022615           Sequence 8286         AK001392         Genbank         7022621           Sequence 8287         AK001395         Genbank         7022625           Sequence 8288         AK001410         Genbank         7022649           Sequence 8289         AK001436         Genbank         7022692           Sequence 8290         AK001478         Genbank         7022760           Sequence 8291         AK001492         Genbank         7022781           Sequence 8292         AK001508         Genbank         7022805           Sequence 8293         AK001521         Genbank         7022828				
Sequence 8284         AK001366         Genbank         7022579           Sequence 8285         AK001387         Genbank         7022615           Sequence 8286         AK001392         Genbank         7022621           Sequence 8287         AK001395         Genbank         7022625           Sequence 8288         AK001410         Genbank         7022649           Sequence 8289         AK001436         Genbank         7022692           Sequence 8290         AK001478         Genbank         7022760           Sequence 8291         AK001492         Genbank         7022781           Sequence 8292         AK001508         Genbank         7022805           Sequence 8293         AK001521         Genbank         7022828			Genbank	
Sequence 8285         AK001387         Genbank         7022615           Sequence 8286         AK001392         Genbank         7022621           Sequence 8287         AK001395         Genbank         7022625           Sequence 8288         AK001410         Genbank         7022649           Sequence 8289         AK001436         Genbank         7022692           Sequence 8290         AK001478         Genbank         7022760           Sequence 8291         AK001492         Genbank         7022781           Sequence 8292         AK001508         Genbank         7022805           Sequence 8293         AK001521         Genbank         7022828			Genbank	
Sequence 8286         AK001392         Genbank         7022621           Sequence 8287         AK001395         Genbank         7022625           Sequence 8288         AK001410         Genbank         7022649           Sequence 8289         AK001436         Genbank         7022692           Sequence 8290         AK001478         Genbank         7022760           Sequence 8291         AK001492         Genbank         7022781           Sequence 8292         AK001508         Genbank         7022805           Sequence 8293         AK001521         Genbank         7022828			Genbank	
Sequence 8287         AK001395         Genbank         7022625           Sequence 8288         AK001410         Genbank         7022649           Sequence 8289         AK001436         Genbank         7022692           Sequence 8290         AK001478         Genbank         7022760           Sequence 8291         AK001492         Genbank         7022781           Sequence 8292         AK001508         Genbank         7022805           Sequence 8293         AK001521         Genbank         7022828				
Sequence 8288         AK001410         Genbank         7022649           Sequence 8289         AK001436         Genbank         7022692           Sequence 8290         AK001478         Genbank         7022760           Sequence 8291         AK001492         Genbank         7022781           Sequence 8292         AK001508         Genbank         7022805           Sequence 8293         AK001521         Genbank         7022828	•		Genbank	
Sequence 8289         AK001436         Genbank         7022692           Sequence 8290         AK001478         Genbank         7022760           Sequence 8291         AK001492         Genbank         7022781           Sequence 8292         AK001508         Genbank         7022805           Sequence 8293         AK001521         Genbank         7022828				
Sequence 8290         AK001478         Genbank         7022760           Sequence 8291         AK001492         Genbank         7022781           Sequence 8292         AK001508         Genbank         7022805           Sequence 8293         AK001521         Genbank         7022828			Genbank	
Sequence 8291         AK001492         Genbank         7022781           Sequence 8292         AK001508         Genbank         7022805           Sequence 8293         AK001521         Genbank         7022828	•		Genbank	
Sequence 8292         AK001508         Genbank         7022805           Sequence 8293         AK001521         Genbank         7022828		AK001478	Genbank	
Sequence 8293 AK001521 Genbank 7022828	Sequence 8291		Genbank	
• • • • • • • • • • • • • • • • • • • •	Sequence 8292		Genbank	7022805
Sequence 8294 AK001539 Genbank 7022856	Sequence 8293	AK001521	Genbank	7022828
	Sequence 8294	AK001539	Genbank	7022856

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Sequence 8295	AK001565	Genbank	7022895
Sequence 8296	AK001601	Genbank	7022955
Sequence 8297	AK001613	Genbank	7022975
Sequence 8298	AK001627	Genbank	7022996
Sequence 8299	AK001636	Genbank	7023010
Sequence 8300	AK001675	Genbank	7023079
Sequence 8301	AK001690	Genbank	7023105
	AK001691	Genbank	7023106
Sequence 8302		Genbank	7023122
Sequence 8303	AK001700		7023122
Sequence 8304	AK001714	Genbank	
Sequence 8305	AK001716	Genbank	7023150
Sequence 8306	AK001725	Genbank	7023165
Sequence 8307	AK001741	Genbank	7023192
Sequence 8308	AK001775	Genbank	7023255
Sequence 8309	AK001829	Genbank	7023339
Sequence 8310	AK001852	Genbank	7023376
Sequence 8311	AK001939	Genbank	7023513
Sequence 8312	AK001962	Genbank	7023551
Sequence 8313	AK002019	Genbank	7023647
Sequence 8314	AK002051	Genbank	7023698
Sequence 8315	AK002060	Genbank	7023713
Sequence 8316	AK002094	Genbank	7023767
Sequence 8317	AK002136	Genbank	7023830
Sequence 8318	AK002165	Genbank	7023876
Sequence 8319	AK002212	Genbank	7023953
Sequence 8320	AL008639	Genbank	6599068
Sequence 8321	AL008721	Genbank	3171883
Sequence 8322	AL020995	Genbank	5881341
Sequence 8323	AL021453	Genbank	3413288
Sequence 8324	AL021546	Genbank	2826890
Sequence 8325	AL021707	Genbank	4582132
Sequence 8326	AL021807	Genbank	9367202
Sequence 8327	AL021917	Genbank	9367203
Sequence 8328	AL021978	Genbank	3135963
Sequence 8329	AL022157	Genbank	3319676
Sequence 8330	AL022313	Genbank	4200326
Sequence 8331	AL022318	Genbank	4826439
Sequence 8332	AL022339	Genbank	3355600
Sequence 8333	AL022394	Genbank	9581758
Sequence 8334	AL023803	Genbank	9408726
Sequence 8335	AL024498	Genbank	5924006
Sequence 8336	AL031387	Genbank	4493578
Sequence 8337	AL031307	Genbank	4835258
Sequence 8338	AL031680	Genbank	9368417
Sequence 8339	AL031685	Genbank	9368423
Sequence 6339	AL031733	Genbank	4826495
Sequence 8340	AL031733 AL031778	Genbank	4153958
Sequence 8341	AL034743	Genbank	5931807
Sequence 8342			5918013
Sequence 8343	AL034379	Genbank Genbank	7981257
Sequence 8344	AL034380		591816 <b>1</b>
Sequence 8345	AL035420	Genbank	4826429
Sequence 8346	AL035551	Genbank	6002306
Sequence 8347	AL035587	Genbank	6562059
Sequence 8348	AL035659	Genbank	
Sequence 8349	AL035683	Genbank	7288039 4499949
Sequence 8350	AL049219	Genbank	<del></del>

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Sequence 8351	AL049367	Genbank	4500158
Sequence 8352	AL049539	Genbank	6456855
Sequence 3353	AL049546	Genbank	4678775
Sequence 8354	AL049589	Genbank	
Sequence 8355	AL049611	Genbank	5679 <b>5</b> 67
Sequence 8356	AL049646		5881343
Sequence 8357	AL049712	Genbank	9368462
	- · · · · -	Genbank	5629919
Sequence 8358	AL049713	Genbank	6006528
Secuence 8359	AL049742	Genbank	5419783
Secuence 8360	AL049795	Genbank	6010175
Secuence 8361	AL049801	Genbank	4902677
Secuence 8362	AL049821	Genbank	5791513
Sequence 8363	AL0499 <b>4</b> 3	Genbank	4884187
Sequence 8364	AL050011	Genbank	4884080
Sequence 8365	AL050028	Genbank	4884267
Sequence 8366	AL050107	Genbank	4884135
Sequence 8367	AL050118	Genbank	4884143
Sequence 8368	AL050135	Genbank	4884344
Sequence 8369	AL050187	Genbank	4884402
Sequence 8370	AL050208	Genbank	4884447
Sequence 8371	AL050274	Genbank	4886452
Sequence 8372	AL050332	Genbank	6010176
Sequence 8373	AL050367	Genbank	4914600
Sequence 8374	AL050396	Genbank	4914618
Sequence 8375	AL078581	Genbank	6967288
Sequence 8376	AL078594	Genbank	7161750
Sequence 8377	AL078611	Genbank	5019312
Sequence 8378	AL078621	Genbank	6013067
Sequence 8379	AL079298	Genbank	5102736
Sequence 8380	AL079352	Genbank	7799779
Sequence 8381	AL080080	Genbank	5262491
Sequence 8382	AL080155	Genbank	5262612
Sequence 8383	AL080157	Genbank	5262616
Sequence 8384	AL080172	Genbank	5262642
Sequence 8385	AL096738	Genbank	5419870
Sequence 8386	AL096740	Genbank	5419896
Sequence 8387	AL096752	Genbank	5419888
Sequence 8388	AL096791	Genbank	6066332
Sequence 8389	AL109788	Genbank	5725476
Sequence 8390	AL109794	Genbank	5725481
Sequence 8391	AL109925	Genbank	7105935
Sequence 8392	AL110244	Genbank	5817191
Sequence 8393	AL110268	Genbank	5817041
Sequence 8394	AL110273	Genbank	5817091
Sequence 8395	AL117398	Genbank	5911918
Sequence 8396	AL117409	Genbank	5912046
Sequence 8397	AL117429	Genbank	5911863
Sequence 8398	AL117482	Genbank	5911960
Sequence 8399	AL117500	Genbank	5912005
Sequence 8400	AL117572	Genbank	5912124
Sequence 8401	AL117573	Genbank	5912125
Sequence 8402	AL117579	Genbank	5912135
Secuence 8403	AL117594	Genbank	5912158
Sequence 8404	AL117607	Genbank	5912177
Sequence 8405	AL117694	Genbank	8248721
Secuence 8406	AL121575	Genbank	7264053
		CHIDAIR	1204003

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Sequence 8407	AL121790	Genbank	8919824
Sequence 8408	AL122020	Genbank	8176920
Sequence 8409	AL133C36	Genbank	6453441
Sequence 8410	AL133064	Genbank	6453490
	AL133078	Genbank	6453522
Sequence 8411			7228177
Sequence 8412	AL133215	Genbank	
Sequence 8413	AL133246	Genbank	7594587
Sequence 8414	AL133353	Genbank	6706037
Sequence 8415	AL133455	Genbank	7019734
Sequence 8416	AL133555	Genbank	6599122
Sequence 8417	AL133583	Genbank	6599167
Sequence 8418	AL133589	Genbank	6599178
Sequence 8419	AL133609	Genbank	6599%
Sequence 8420	AL135744	Genbank	6682
Sequence 8421	AL135858	Genbank	715962
Sequence 8422	AL136295	Genbank	<b>685093</b> ର
Sequence 8423	AL136543	Genbank	6 <b>80764</b> 6
Sequence 8424	AL137263	Genbank	6807 <b>69</b> 2
Sequence 8425	AL137304	Genbank	68077 <b>68</b>
Sequence 8426	AL137327	Genbank	6807 <b>818</b>
Sequence 8427	AL137406	Genbank	6807955
Sequence 8428	AL137471	Genbank	6808079
Sequence 8429	AL137540	Genbank	6808218
Sequence 8430	AL137548	Genbank	6808231
Sequence 8431	AL137585	Genbank	6808304
Sequence 8432	AL137641	Genbank	6808443
Sequence 8433	AL137681	Genbank	6807931
Sequence 8434	AL137683	Genbank	6807935
Sequence 8435	AL137727	Genbank	6808256
Sequence 8436	AL137743	Genbank	6808323
Sequence 8437	AL139165	Genbank	6969146
Sequence 8438	AP000010	Genbank	4666256
Sequence 8430	AP000016	Genbank	3132356
Sequence 8439	AP000046	Genbank	4579987
Sequence 8440	AP000493	Genbank	5926660
Sequence 8441	AP000493 AP000501	Genbank	5926688
Sequence 8442	AP000529	Genbank	5931507
Sequence 8443	AP000529 AP000694	Genbank	6693626
Sequence 8444		Genbank	6899854
Sequence 8445	AP001115	Genbank	6970361
Sequence 8446	AP001137	Genbank	220051
Sequence 8447	D00591	Genbank	220140
Sequence 8448	D01038	Genbank	219899
Sequence 8449	D10040		520586
Sequence 8450	D10495 D12981	Genbank Genbank	219979
Sequence 8451	= .=		285909
Sequence 8452	D13119	Genbank	496368
Sequence 8453	D13286	Genbank	220071
Sequence 8454	D13540	Genbank	286010
Sequence 8455	D13627	Genbank	
Sequence 8456	D13635	Genbank	285982
Sequence 8457	D13748	Genbank	219402
Sequence 8458	D14662	Genbank	285948
Sequence 8459	D14689	Genbank	285956
Sequence 8460	D14694	Genbank	603801
Sequence 8461	D14811	Genbank	285966
Sequence 8462	D16307	Genbank	30 <b>3594</b>

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Sequence 8463	D16562	Genbank	506336
Sequence 8464	D21063	Genbank	434752
Sequence 8465	D21153	Genbank	434758
Sequence 8466	D21235	Genbank	498145
Sequence 8467	D21254	Genbank	575577
Sequence 8468	D21261	Genbank	434762
Sequence 8469	D21262	Genbank	434764
Sequence 8470	D21852	Genbank	434768
Sequence 8471	D25215	Genbank	517114
Sequence 8472	D26156	Genbank	505087
Sequence 8473	D26181	Genbank	473712
Sequence 8474	D26351	Genbank	450470
Sequence 8475	D26485	Genbank	468934
Sequence 8476	D28473	Genbank	551621
Sequence 8477	D28476	Genbank	460710
Sequence 8478	D28589	Genbank	460714
Sequence 8479	D29954	Genbank	473940
Sequence 8480	D29958	Genbank	473948
Sequence 8481	D30756	Genbank	488500
Sequence 8482	D31762	Genbank	498149
Sequence 8483	D31764	Genbank	498153
Sequence 8484	D317 <b>67</b>	Genbank	505091
Sequence 8485	D31885	Genbank	505097
Sequence 8486	D31888	Genbank	506340
Sequence 8487	D37766	Genbank	2429078
Sequence 8488	D38047	Genbank	1037163
Sequence 8489	D38048	Genbank	1531532
Sequence 8490	D38073	Genbank	862331
Sequence 8491	D38522	Genbank	559331
Sequence 8492	D38524	Genbank	633070
Sequence 8493	D38549	Genbank	559702
Sequence 8494	D38553	Genbank	55 <b>9</b> 714
Sequence 8495	D38555	Genbank	559716
Sequence 8496	D38616	Genbank	1304117
Sequence 8497	D42047	Genbank	577306
Sequence 8498	D42055	Genbank	577312
Sequence 8499 Sequence 8500	D42085	Genbank	577316
Sequence 8501	D43682 D43949	Genbank	1060913
Sequence 8502	D43949 D43950	Genbank	603952
Sequence 8503	D43950 D43951	Genbank Genbank	603954
Sequence 8504	D45931 D45021	Genbank Genbank	603956
Sequence 8505	D45021 D45131	Genbank Genbank	624872
Sequence 8506	D49489	Genbank	1304103 1136742
Sequence 8507	D49499 D49490	Genbank	1072306
Sequence 8508	D50369		
Sequence 8509	D50309	Genbank Genbank	2605589
Sequence 8510	D50923	Genbank	2605593
Sequence 8511	D50929	Genbank	1469188 1469200
Sequence 8512	D55716	Genbank	
Sequence 8513	D63475	Genbank	1255616 1665724
Sequence 8514	D63478	Genbank	1469869
Sequence 8515	D63479	Genbank	6633998
Sequence 8516	D63486	Genbank	1469885
Sequence 8517	D63780	Genbank	2196444
Sequence 8518	D63861	Genbank	1769811
	20001	CCIDAIR	1705011

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Sequence 8519	D63875	Genbank	961441
Sequence 8520	D78014	Genbank	1330241
•	D79205	Genbank	1754620
Sequence 8521			
Sequence 8522	D79984	Genbank	1136385
Sequence 8523	D79991	Genbank	1136397
Sequence 8524	D80000	Genbank	1136415
Sequence 8525	D80009	Genbank	1136433
•	D80010	Genbank	1136435
Sequence 8526			1311461
Sequence 8527	D82348	Genbank	
Sequence 8528	D83077	Genbank	1304131
Sequence 8529	D83778	Genbank	1228038
Seguence 8530	D83781	Genbank	1228044
Sequence 8531	D83782	Genbank	1228046
Sequence 8532	D84064	Genbank	2618587
	D86198	Genbank	3062805
Sequence 8533			1503997
Sequence 8534	D86962	Genbank	
Sequence 8535	D86965	Genbank	1504003
Sequence 8536	D86977	Genbank	1504027
Sequence 8537	D86978	Genbank	1504029
Sequence 8538	D86985	Genbank	6634002
Sequence 8539	D86987	Genbank	3869186
	D87127	Genbank	1817 <b>551</b>
Sequence 8540			
Sequence 8541	D87436	Genbank	1665766
Sequence 8542	D87447	Genbank	1665782
Sequence 8543	D87452	Genbank	1665792
Sequence 8544	D87464	Genbank	1665812
Sequence 8545	D87470	Genbank	1665822
Sequence 8546	D87717	Genbank	1663709
Sequence 8547	D87735	Genbank	1620021
	D88674	Genbank	2641951
Sequence 8548		Genbank	1843433
Sequence 8549	D88687		
Sequence 8550	D89980	Genbank	2804272
Sequence 8551	D90391	Genbank	219491
Sequence 8552	E01094	Genbank	21 <b>69353</b>
Sequence 8553	E01787	Genbank	2170040
Sequence 8554	E01816	Genbank	2170068
Sequence 8555	E01956	Genbank	2170204
Sequence 8556	E02164	Genbank	2170402
	E02628	Genbank	2170856
Sequence 8557			2174903
Sequence 8558	E06721	Genbank	
Sequence 8559	E12458	Genbank	3251291
Sequence 8560	E13330	Genbank	3252135
Sequence 8561	E14580	Genbank	57 <b>09263</b>
Sequence 8562	E14872	Genbank	57 <b>09555</b>
Sequence 8563	E15239	Genbank	5709922
Sequence 8564	E15695	Genbank	5710378
Sequence 8565	E17361	Genbank	5712044
		Genbank	182862
Sequence 8566	J02642		
Sequence 8567	J02923	Genbank	189501
Sequence 8568	J03007	Genbank	179211
Sequence 8569	J03015	Genbank	33 <b>7755</b>
Sequence 8570	J03202	Genbank	1 <b>86916</b>
Sequence 8571	J03473	Genbank	337423
Sequence 8572	J03518	Genbank	340389
Sequence 8573	J03518	Genbank	188671
•	J03526 J03575	Genbank	189737
Sequence 8574	103575	Genbank	103131

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Sequence 857	5 J03576	Genbank	189753
Sequence 857	6 J03592	Genbank	339722
Sequence 857	7 J03799	Genbank	186840
Sequence 857	8 J03827	Genbank	340418
Sequence 857		Genbank	187464
Sequence 858		Genbank	292829
Sequence 858		Genbank	177801
Sequence 858		Genbank	551173
Sequence 858		Genbank	181239
Sequence 858		Genbank	338243
Sequence 858			
Sequence 858		Genbank	178649
•		Genbank	338246
Sequence 858		Genbank	184422
Sequence 858		Genbank	1203981
Sequence 858		Genbank	186504
Sequence 859		Genbank	340020
Sequence 859		Genbank	189904
Sequence 859		Genbank	292032
Sequence 859		Genbank	179282
Sequence 859		Genbank	307084
Sequence 859	5 L05093	Genbank	401844
Sequence 859	6 L05425	Genbank	179284
Sequence 859	7 L06237	Genbank	437000
Sequence 859	8 L06328	Genbank	340200
Sequence 8599		Genbank	190073
Sequence 860		Genbank	291894
Sequence 860		Genbank	307198
Sequence 860	2 L09159	Genbank	307374
Sequence 860		Genbank	3192937
Sequence 860		Genbank	307421
Sequence 860		Genbank	307423
Sequence 860		Genbank	292165
Sequence 860		Genbank	307437
Sequence 860		Genbank	307355
Sequence 860		Genbank	348238
Sequence 861		Genbank	292937
Sequence 861		Genbank	452438
Sequence 861		Genbank	306467
Sequence 861		Genbank	306456
Sequence 861		Genbank	
Sequence 861			507251
		Genbank	388308
Sequence 8610		Genbank	347313
Sequence 861		Genbank	409222
Sequence 8618		Genbank	414316
Sequence 8619		Genbank	457136
Sequence 862		Genbank	414584
Sequence 862		Genbank	450272
Sequence 862		Genbank	457232
Sequence 862:		Genbank	<b>558656</b>
Sequence 862		Genbank	<b>80</b> 80 <b>06</b>
Sequence 862		Genbank	632965
Sequence 8620	6 L29296	Genbank	559042
Sequence 862	7 L29555	Genbank	522196
Sequence 862	8 L34075	Genbank	508481
Secuence 8629		Genbank	561631
Secuence 8636	D L36818	Genbank	556191

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Sequence 8631	L38961	Genbank	624703
Sequence 8632	L39068	Genbank	994714
Sequence 8633	L39210	Genbank	1702963
Sequence 8634	L39211	Genbank	755645
Sequence 8635	L40326	Genbank	695361
Sequence 8636	L40586	Genbank	1222629
Sequence 8637	L40933	Genbank	1160964
Sequence 8638	L41607	Genbank	886293
Sequence 8639	L41816	Genbank	790789
	L41319	Genbank	1203964
Sequence 8640	L42579	Genbank	974142
Sequence 8641	L42542 L42809	Genbank	1160966
Sequence 8642			896282
Sequence 8643	L43509	Genbank	1082035
Sequence 8644	L47162	Genbank	
Sequence 8645	L47647	Genbank	1000861
Sequence 8646	L49345	Genbank	1405422
Sequence 8647	L54057	Genbank	1196416
Sequence 8648	L77701	Genbank	1280205
Sequence 8649	L78464	Genbank	1479980
Sequence 8650	L78669	Genbank	1374862
Sequence 8651	M11147	Genbank	182513
Sequence 8652	M11560	Genbank	178350
Sequence 8653	M12938	Genbank	182515
Sequence 8654	M13520	Genbank	<sub>/</sub> 179459
Sequence 8655	M13755	Genbank	184570
Sequence 8656	M14326	Genbank	340005
Sequence 8657	M14631	Genbank	183416
Sequence 8658	M15182	Genbank	183232
Sequence 8659	M16038	Genbank	187268
Sequence 8660	M16247	Genbank	1780 <b>44</b>
Sequence 8661	M16279	Genbank	188542
Sequence 8662	M17081	Genbank	178151
Sequence 8663	M17851	Genbank	182860
Sequence 8664	M17885	Genbank	190231
Sequence 8665	M17886	Genbank	190233
Sequence 8666	M18112	Genbank	190 <b>166</b>
Sequence 8667	M18157	Genbank	186640
Sequence 8668	M18366	Genbank	179131
Sequence 8669	M19961	Genbank	180940
Sequence 8670	M20372	Genbank	189372
Sequence 8671	M20776	Genbank	180909
Sequence 8672	M21154	Genbank	17 <b>8517</b>
Sequence 8673	M21575	Genbank	1311702
Sequence 8674	M21895	Genbank	189523
Sequence 8675	M22382	Genbank	190126
Sequence 8676	M22538	Genbank	986883
Sequence 8677	M22803	Genbank	190379
Sequence 8678	M22919	Genbank	189016
Sequence 8679	M22920	Genbank	189021
Sequence 8680	M22920	Genbank	190282
Sequence 8681	M23254	Genbank	511636
Sequence 96001	M23419	Genbank	181996
Sequence 8682	M23725	Genbank	189997
Sequence 8683	M24194	Genbank Genbank	187701
Sequence 8684		Genbank Genbank	182604
Sequence 8685	M24355	Genbank Genbank	190785
Sequence 8686	M24486	Genbank	130703

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Sequence 8	3687	M24543	Genbank	341200
Sequence 8	3688	M26066	Genbank	189854
Sequence 8		M26325	Genbank	186688
Sequence 8		M26663	Genbank	618463
Sequence 8		M29366	Genbank	181979
•		M29551	Genbank	
Sequence 8				180708
Sequence 8		M31159	Genbank	183115
Sequence 8		M31511	Genbank	179907
Sequence 8		M31606	Genbank	189940
Sequence 8		M33308	Genbank	340236
Sequence 8	3697	M34055	Genbank	190791
Sequence 8	3698	M34064	Genbank	416292
Sequence 8	3699	M34081	Genbank	337458
Sequence 8		M34088	Genbank	182122
Sequence 8		M34668	Genbank	190738
Sequence 8		M37499	Genbank	188886
Sequence 8		M38188	Genbank	189378
Sequence 8		M55153	Genbank	339520
•		M55618		
Sequence 8			Genbank	184483
Sequence 8		M57567	Genbank	178986
Sequence 8		M58028	Genbank	340071
Sequence 8		M59371	Genbank	181943
Sequence 8		M59849	Genbank	182591
Sequence 8		M61715	Genbank	<b>34</b> 0367
Sequence 8		M62810	Genbank	188563
Sequence 8		M64098	Genbank	183891
Sequence 8	3713	M64571	Genbank	187382
Sequence 8	3714	M64788	Genbank	190855
Sequence 8	3715	M65217	Genbank	184404
Sequence 8	3716	M67480	Genbank	190364
Sequence 8		M69066	Genbank	188625
Sequence 8		M69181	Genbank	641957
Sequence 8		M73816	Genbank	183609
Sequence 8		M74491	Genbank	178161
Sequence 8		M75126	Genbank	184020
Sequence 8		M76729	Genbank	189519
Sequence 8		M80254	Genbank	181273
•			Genbank	
Sequence 8		M80335	* *	189966
Sequence 8		M80646	Genbank	338701
Sequence 8		M80776	Genbank	179334
Sequence 8		M81934	Genbank	180172
Sequence 8		M83651	Genbank	431032
Sequence 8		M84326	Genbank	178163
Sequence 8		M84739	Genbank	179881
Sequence 8		M85038	Genbank	190189
Sequence 8	3732	M86492	Genbank	18336 <b>9</b>
Sequence 8	3733	M91670	Genbank	181915
Sequence 8	3734	M95627	Genbank	870802
Sequence 8		M96684	Genbank	190749
Sequence 8		M96803	Genbank	338442
Sequence 8		M96995	Genbank	181975
Sequence 8		M97501	Genbank	180621
Sequence 8		M97934	Genbank	186566
Sequence 8		M97935	Genbank	2281070
Sequence 8		M98478		
•			Genbank	339577
Sequence 8	142	M99422	Genbank	183662

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Sequence 8743	S60710	Genbank	300301
Sequence 8744	S62138	Genbank	386158
Sequence 8745	S67779	Genbank	459372
Sequence 8746	S69272	Genbank	546087
Sequence 8747	S70114	Genbank	546696
Sequence 8748	S73498	Genbank	688010
Sequence 8749	S74678	Genbank	241477
	-		
Sequence 8750	S74799	Genbank	797365
Sequence 8751	S75174	Genbank	802120
Sequence 8752	S82081	Genbank	1488412
Sequence 8753	U00930	Genbank	405043
Sequence 8754	U00947	Genbank	405049
Sequence 8755	U02390	Genbank	409928
Sequence 8756	U02493	Genbank	407307
Sequence 8757	U03886	Genbank	458225
Sequence 8758	U05572	Genbank	1419373
Sequence 8759	U07563	Genbank	514264
Sequence 8760	U07681	Genbank	706838
Sequence 8761	U07810	Genbank	469038
Sequence 8762	U07919	Genbank	995897
Sequence 8763	U08377	Genbank	508230
Sequence 8764	U09220	Genbank	799338
Sequence 8765	U09284	Genbank	516011
Sequence 8766	U09716	Genbank	606827
•	U09813	Genbank	1008454
Sequence 8767			984280
Sequence 8768	U10248	Genbank	
Sequence 8769	U10323	Genbank	532312
Sequence 8770	U10439	Genbank	577169
Sequence 8771	U10691	Genbank	533522
Sequence 8772	U10886	Genbank	1685074
Sequence 8773	U12424	Genbank	525319
Sequence 8774	U12535	Genbank	530822
Sequence 8775	U13022	Genbank	537293
Sequence 8776	U13991	Genbank	562076
Sequence 8777	U14193	Genbank	555899
Sequence 8778	U14966	Genbank	550012
Sequence 8779	U14969	Genbank	550018
Sequence 8780	U14970	Genbank	5 <b>50</b> 020
Sequence 8781	U14971	Genbank	5 <b>50022</b>
Sequence 8782	U15158	Genbank	576622
Sequence 8783	U16660	Genbank	5640 <b>6</b> 4
Sequence 8784	U18121	Genbank	915283
Sequence 8785	U18197	Genbank	603073
Sequence 8786	U18937	Genbank	899108
Sequence 8787	U19145	Genbank	914904
Sequence 8788	U19252	Genbank	808035
Sequence 8789	U20476	Genbank	3077819
Sequence 8790	U21910	Genbank	736402
Sequence 8791	U22055	Genbank	799176
Sequence 8792	U22815	Genbank	930340
Sequence 8793	U23028	Genbank	806853
Sequence 8794	U24169	Genbank	1215668
Sequence 8795	U24223	Genbank	1215670
Sequence 8796	U25816	Genbank	837262
Sequence 8797	U26425	Genbank	836 <b>66</b> 4
Sequence 8798	U27515	Genbank	4096274
ocquerioe or 30	02/3/3	Genbank	4030Z/4

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Sequence 8799	U28337	Genbank	881950
Sequence 8800	U28635	Genbank	881953
Sequence 8801	U28727	Genbank	1142969
Sequence 8802	U28811	Genbank	
Sequence 8803	U30255	Genbank	1373018
Sequence 8804	U32944	Genbank	984324
Sequence 8805			1209060
	U32939	Genbank	993045
Sequence 8806	U33460	Genbank	4096590
Sequence 8807	U33760	Genbank	995823
Sequence 8808	U33818	Genbank	1163176
Sequence 8809	U34038	Genbank	1041728
Sequence 8810	U34877	Genbank	1143231
Sequence 8811	U34994	Genbank	6525311
Sequence 8812	U35116	Genbank	1008541
Sequence 8813	U36138	Genbank	1244507
Sequence 8814	U36764	Genbank	1036804
Sequence 8815	U37426	Genbank	1171152
Sequence 8816	U37558	Genbank	1017812
Sequence 8817	U38784	Genbank	1574947
Sequence 8818	U38894	Genbank	1589739
Sequence 8819	U38945	Genbank	1353569
Sequence 8820	U39400	Genbank	1234796
Sequence 8821	U39945	Genbank	1209686
Sequence 8822	U40152	Genbank	1113100
Sequence 8823	U40989	Genbank	1125005
Sequence 8824	U41371	Genbank	1173904
Sequence 8825	U41387	Genbank	
Sequence 8826	U41806	Genbank	1230563
Sequence 8827	U46691	Genbank	1145798
Sequence 8828	U47742	Genbank	1401077
Sequence 8829	U49436	Genbank	1517913
Sequence 8830	U50079	Genbank	1229139
Sequence 8831	U51587	Genbank	1277083
Sequence 8832	U51625		1669823
Sequence 8833	U51678	Genbank	1695798
Sequence 8834	U53209	Genbank	1915966
Sequence 8835	U53346	Genbank	1256836
Sequence 8836		Genbank	1545981
Sequence 8837	U53530 U54559	Genbank	1314642
Sequence 8838		Genbank	2351379
Sequence 8839	U <b>5</b> 7052	Genbank	1519039
Sequence 8840	U58766	Genbank	1381178
Sequence 8841	U59435	Genbank	2697004
	U61262	Genbank	1621606
Sequence 8842	U61397	Genbank	1518693
Sequence 8843	U61734	Genbank	1407825
Sequence 8844	U61843	Genbank	3650451
Sequence 8845	U62317	Genbank	68 <b>62558</b>
Sequence 8846	U62583	Genbank	1778 <b>05</b> 0
Sequence 8847	U63296	Genbank	2047312
Sequence 8848	U63743	Genbank	1695881
Sequence 8849	U64791	Genbank	1498734
Sequence 8850	U65676	Genbank	1654350
Sequence 8851	U65928	Genbank	1549382
Sequence 8852	U66618	Genbank	1549244
Sequence 8853	U67058	Genbank	4097702
Sequence 8854	U68723	Genbank	2114391

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Sequence 8855	U70063	Genbank	1743866
Sequence 8856	U70312	Genbank	2 <b>8652</b> 18
Sequence 8857	U70451	Genbank	1763090
Sequence 8858	U72069	Genbank	1657775
Sequence 8859	U72398	Genbank	1622940
Sequence 8860	U72515	Genbank	1673519
Sequence 8861	U73167	Genbank	2880032
Sequence 8862	U73338	Genbank	1763268
	U74612	Genbank	1842252
Sequence 8863			
Sequence 8864	U74613	Genbank	1842254
Sequence 8865	U75329	Genbank	2507612
Sequence 8866	U75330	Genbank	2507614
Sequence 8867	U76638	Genbank	1710174
Sequence 8868	U76764	Genbank	1685050
Sequence 8869	U77456	Genbank	1679778
		Genbank	
Sequence 8870	U79241		1710187
Sequence 8871	U79302	Genbank	1710287
Sequence 8872	U79458	Genbank	4205085
Sequence 8873	U80040	Genbank	1718501
Sequence 8874	U80735	Genbank	2565045
Sequence 8875	U80811	Genbank	1857424
Sequence 8876	U81002	Genbank	4580010
		-	1764089
Sequence 8877	U81599	Genbank	
Sequence 8878	U82226	Genbank	1848270
Sequence 8879	U82535	Genbank	2149155
Sequence 8880	U83410	Genbank	1923242
Sequence 8881	U856 <b>58</b>	Genbank	2058552
Sequence 8882	U89278	Genbank	1877500
Sequence 8883	U89505	Genbank	2078528
Sequence 8884	U89995	Genbank	2078532
•	U90339	Genbank	1906010
Sequence 8885		Genbank	2290529
Sequence 8886	U94747		
Sequence 8887	U94788	Genbank	3041866
Sequence 8888	U96876	Genbank	2358268
Sequence 8889	U97156	Genbank	2459796
Sequence 8890	U97519	Genbank	2213812
Sequence 8891	V00572	Genbank	35434
Sequence 8892	X01060	Genbank	37432
Sequence 8893	X02152	Genbank	34312
Sequence 8894	X03363	Genbank	31197
	X03473	Genbank	32106
Sequence 8895			
Sequence 8896	X04408	Genbank	31914
Sequence 8897	X04526	Genbank	31667
Sequence 8898	X04741	Genbank	35439
Sequence 8899	X05332	Genbank	35740
Sequence 8900	X06272	Genbank	30865
Sequence 8901	X06409	Genbank	30204
Sequence 8902	X06562	Genbank	31737
Sequence 8903	X06747	Genbank	36101
Sequence 8904	X06994	Genbank	30302
Sequence 8905	X12433	Genbank	32451
Sequence 8906	X12597	Genbank	32326
Sequence 8907	X12660	Genbank	36654
Sequence 8908	X13294	Genbank	28685
Sequence 8909	X13916	Genbank	34338
Sequence 8910	X13939	Genbank	34237

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Sequence 8911	X15488	Genbank	30933
Sequence 8912	X15729	Genbank	38317
Sequence 8913	X158 <b>80</b>	Genbank	30029
Sequence 8914	X16 <b>064</b>	Genbank	37495
Sequence 8915	X16135	Genbank	32355
Sequence 8916	X16396	Genbank	35070
Sequence 8917	X17567	Genbank	36512
Sequence 8918	X17620	Genbank	35067
Eguence 8919	X51466	Genbank	31105
Sequence 8920	X52519	Genbank	36694
Sequence 8921	X52851	Genbank	30167
Sequence 8922	X52955	Genbank	29627
Sequence 8923	X532 <b>8</b> 0		
Sequence 8924		Genbank	29504
	X56998	Genbank	37564
Sequence 8925	X57129	Genbank	31967
Sequence 8926	X57206	Genbank	33990
Sequence 8927	X59618	Genbank	36154
Sequence 8928	X61123	Genbank	29508
Sequence 8929	X64123	Genbank	35819
Sequence 8930	X64229	Genbank	30502
Sequence 8931	X64707	Genbank	29382
Sequence 8932	X64875	Genbank	398163
Sequence 8933	X66 <b>364</b>	Genbank	36620
Sequence 8934	X67155	Genbank	6 <b>723674</b>
Sequence 8935	X67334	Genbank	468703
Sequence 8936	X68 <b>836</b>	Genbank	36326
Sequence 8937	X70940	Genbank	38455
Sequence 8938	X71 <b>428</b>	Genbank	393415
Sequence 8939	X71 <b>81</b> 0	Genbank	297905
Sequence 8940	X74070	Genbank	395086
Sequence 8941	X74794	Genbank	683749
Sequence 8942	X76057	Genbank	416016
Sequence 8943	X78136	Genbank	460772
Sequence 8944	X78706	Genbank	471311
Sequence 8945	X80197	Genbank	951271
Sequence 8946	X80230	Genbank	599828
Sequence 8947	X80910	Genbank	531475
Sequence 8948	X81197	Genbank	773572
Sequence 8949	X81889	Genbank	1702923
Sequence 8950	X82321	Genbank	1617117
Sequence 8951	X85137	Genbank	1155083
Sequence 8952	X86012	Genbank	976082
Sequence 8953	X92814	Genbank	1054751
Sequence 8954	X94333	Genbank	1518268
Sequence 8955	X947 <b>54</b>	Genbank	1702931
Sequence 8956	X95762	Genbank	2584786
Sequence 8957	X96 <b>698</b>	Genbank	1418766
Sequence 8958	X97064		
Sequence 8959	X97064 X97124	Genbank Genbank	1296663
Sequence 8960			1518264
•	X97335	Genbank	1507823
Sequence 8961	X97674	Genbank	1877214
Sequence 8962	X977 <b>9</b> 5	Genbank	1495482
Sequence 8963	X9 <b>8054</b>	Genbank	1359754
Sequence 8964	X98261	Genbank	1770459
Sequence 8965	X98507	Genbank	1926310
Sequence 8966	X9 <b>9141</b>	Genbank	1903215

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Sequence 8967	X99802	Genbank	2769561
Sequence 8968	Y00052	Genbank	30308
Sequence 8969	Y00281	Genbank	36052
Sequence 8970	Y00282	Genbank	36048
Sequence 8971	Y00815	Genbank	34266
Sequence 8972	Y00978	Genbank	35359
Sequence 8973	Y08134	Genbank	1552274
Sequence 8974	Y09862	Genbank	1945760
Sequence 8975	Y10275	Genbank	1890330
Sequence 8976	Y12478	Genbank	1946204
Sequence 8977	Y13621	Genbank	2190399
Sequence 8978	Y13936	Genbank	2315201
Sequence 8979	Y14486	Genbank	5830437
Sequence 8980	Y15286	Genbank	2584788
Sequence 8981	Y18004	Genbank	4490941
Sequence 8982	Z11583	Genbank	35118
Sequence 8983	Z14244	Genbank	30150
Sequence 8984	Z18956	Genbank	36726
Sequence 8985	Z26649	Genbank	510116
Sequence 8986	Z37166	Genbank	587145
Sequence 8987	Z46376	Genbank	587201
Sequence 8988	Z50022	Genbank	1107702
Sequence 8989	Z58902	Genbank	1030815
Sequence 8990	Z69709	Genbank	1204107
	Z69780	Genbank	1514560
Sequence 8991	Z70200	Genbank	3255964
Sequence 8992	Z71183	Genbank	6580491
Sequence 8993 Sequence 8994	Z71183 Z72499	Genbank	1545951
Sequence 8995	Z73358	Genbank	1322149
Sequence 8996	Z73639	Genbank	2632099
Sequence 8997	Z83840	Genbank	4914518
Sequence 8998	Z84466	Genbank	3319673
Sequence 8999	Z92543	Genbank	3164068
Sequence 9000	Z92545	Genbank	2462414
Sequence 9001	Z93783	Genbank	4581336
Sequence 9002	Z95114	Genbank	5101742
Sequence 9003	Z97053	Genbank	9650676
Sequence 9004	Z98258	Genbank	2462407
Sequence 9005	Z98752	Genbank	9408718
Sequence 9006	Z99716	Genbank	4456457
Sequence 9007	AA011002	EST	1472029
Sequence 9008	AA024935	EST	1489859
Sequence 9009	AA025057	EST	1489962
Sequence 9010	AA025381	EST	1491520
Sequence 9011	AA027341	EST	1492958
Sequence 9012	AA031969	EST	1501922
Sequence 9013	AA033521	EST	1505367
Sequence 9014	AA033724	EST	1505542
Sequence 9015	AA037091	EST	1512209
Sequence 9016	AA037126	EST	1512234
Sequence 9017	AA037364	EST	1512602
Sequence 9018	AA040182	EST	1516598
Sequence 9019	AA043058	EST	1522593
Sequence 9020	AA044122	EST	1521980
Sequence 9021	AA044586	EST	1522977
Sequence 9022	AA044813	EST	1523016
Sequence 9022	~~~~~		323310

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Sequence 9023	AA047046	EST	1524944
Sequence 9024	AA053156	EST	1544427
Sequence 9025	AA053431	EST	1544086
Sequence 9026	AA053679	EST	1544606
Sequence 9027	AA054463	EST	1545388
Sequence 9028	AA055407	EST	1547946
Sequence 9029	AA055649	EST	1547988
Sequence 9030	AA056099	EST	1548436
Sequence 9031	AA056273	EST	1548820
Sequence 9032	AA056358	EST	1548698
Sequence 9033	AA058929	EST	1551784
Sequence 9034	AA062553	EST	1556727
Sequence 9035	AA065298	EST	1929298
Sequence 9036	AA069599	EST	1576958
Sequence 9037	AA070333	EST	1577692
Sequence 9038	AA071122	EST	1578482
Sequence 9039	AA074907	EST	1614776
Sequence 9040	AA075864	EST	1615735
Sequence 9041	AA075873	EST	1615744
Sequence 9042	AA075984	EST	1615853
Sequence 9043	AA077364	EST	1836838
Sequence 9044	AA081041	EST	1622976
Sequence 9045	AA081608	EST	1623666
Sequence 9046	AA082133	EST	1624191
Sequence 9047	AA082281	EST	1624356
Sequence 9048	AA082487	EST	1624560
Sequence 9049	AA082543	EST	1624600
Sequence 9050	AA082817	EST	1624874
Sequence 9051	AA083182	EST	1625258
Sequence 9052	AA083288	EST	1625349
Sequence 9053	AA083300	EST	1625361
Sequence 9054	AA088420	EST	1633967
Sequence 9055	AA098867	EST	1645051
Sequence 9056	AA099415	EST	1645308
Sequence 9057	AA099496	EST	1645449
Sequence 9058	AA099923	EST	1646071
Sequence 9059	AA099926	EST	1646074
Sequence 9060	AA101077	EST	1647597
Sequence 9061	AA101100	EST	1647676
Sequence 9062	AA101277	EST	1648016
Sequence 9063	AA101440 AA111991	EST	1648254
Sequence 9064 Sequence 9065		EST	1664113
Sequence 9066	AA112480	EST	1665157
Sequence 9067	AA112633 AA112679	EST EST	1665334
Sequence 9068			1665397
Sequence 9069	AA115409 AA121091	EST	1670591
Sequence 9070	AA121091 AA121210	EST	1678635
Sequence 9071	AA126105	EST EST	1678949
Sequence 9072	AA126762		1685755
Sequence 9073	AA128709	EST EST	16862 <b>62</b>
Sequence 9074	AA129754	EST	168 <b>86</b> 17 16901 <b>64</b>
Sequence 9075	AA130211	EST	1691215
Sequence 9076	AA130220	EST	1691215
Sequence 9077	AA130247	EST	1691251
Sequence 9078	AA130520	EST	1691803
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Sequence 9079	AA131227	EST	1692735
Sequence 9080	AA131252	EST	1692814
Sequence 9081	AA131474	EST	1692962
Sequence 9082	AA131496	EST	1692984
Sequence 9083	AA132133	EST	1693667
			1693672
Sequence 9084	AA132163	EST	
Sequence 9085	AA132328	EST	1693818
Sequence 9086	AA132414	EST	1693905
Sequence 9087	AA132885	EST	1694372
			1696351
Sequence 9088	AA135266	EST	
Sequence 9089	AA135745	EST	1696757
Sequence 9090	AA135830	EST	1697090
Sequence 9091	AA143151	EST	1712521
			1712672
Sequence 9092	AA143301	EST	
Sequence 9093	AA147806	EST	1717195
Sequence 9094	AA148139	EST	1717530
Sequence 9095	AA149520	EST	1720338
Sequence 9096	AA149637	EST	1720438
Sequence 9097	AA149750	EST	1720892
Sequence 9098	AA150002	EST	1721155
Sequence 9099	AA150076	EST	1721298
Sequence 9100	AA150383	EST	1721914
			1719468
Sequence 9101	AA151213	EST	
Sequence 9102	AA151345	EST	1719890
Sequence 9103	AA151506	EST	1720046
Sequence 9104	AA152172	EST	1721224
Sequence 9105	AA156215	EST	1727895
		EST	1728440
Sequence 9106	AA156807		
Sequence 9107	AA157715	EST	1729340
Sequence 9108	AA158196	EST	1733007
Sequence 9109	AA158600	EST	1733436
Sequence 9110	AA160097	EST	1734545
Sequence 9111	AA160948	EST	1735663
Sequence 9112	AA164729	EST	1740889
Sequence 9113	AA165403	EST	1741436
Sequence 9114	AA165418	EST	1741554
Sequence 9115	AA167191	EST	1745698
Sequence 9116	AA167829	EST	1744980
			1748137
Sequence 9117	AA169752	EST	
Sequence 9118	AA171588	EST	1750657
Sequence 9119	AA174204	EST	1754484
Sequence 9120	AA179201	EST	1760816
Sequence 9121	AA179430	EST	1760782
	AA179442	EST	1760794
Sequence 9122			
Sequence 9123	AA179690	EST	1761923
Sequence 9124	AA181608	EST	1765136
Sequence 9125	AA181618	, EST	1765085
Sequence 9126	AA186685	EST	1774906
		EST	1774847
Sequence 9127	AA186730		
Sequence 9128	AA187424	EST	1773633
Sequence 9129	AA187428	EST	1773747
Sequence 9130	AA187809	EST	1774003
Sequence 9131	AA188497	EST	1775584
•			1775586
Sequence 9132	AA188499	EST	
Sequence 9133	AA188875	EST	1775902
Sequence 9134	AA190915	EST	1779571

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_			
Section 9135	AA191160	EST	17798 <b>54</b>
Secuence 9136	AA194169	EST	1783860
Secuence 9137	AA194896	EST	1784658
Secuence 9138	AA195259	EST	1784959
Sectence 9139	AA195352	EST	1785043
Seclence 9140	AA195510	EST	1785213
Secuence 9141	AA195535	EST	1785248
Secuence 9142	AA196350	EST	1791925
Sectence 9143	AA197018	EST	1792634
Secuence 9144	AA197063	EST	1792629
Sealence 9145	AA197137	EST	1792811
Secuence 9146	AA203450	EST	1799161
Sequence 9147	AA205846	EST	1801217
Secuence 9148	AA205 <b>92</b> 8	EST	1801316
Secuence 9149	AA206583	EST	1801963
Sequence 9150	AA207229	EST	1802920
Sequence 9151	AA209239		
		EST	1807254
Secuence 9152	AA210796	EST	1809450
Sequence 9153	AA211864	EST	1810491
Sequence 9154	AA215319	EST	1815082
Sequence 9155	AA215526	EST	1815298
Sequence 9156	AA215534	EST	1815288
Sequence 9157	AA216769	EST	1817460
Sequence 9158	AA218567	EST	1832642
Sequence 9159	AA218697	EST	1832789
Sequence 9160	AA223382	EST	1843907
Sequence 9161	AA225008	EST	1846317
Sequence 9162	AA226737	EST	1848308
Sequence 9163	AA227227	EST	1848772
Sequence 9164	AA227608	EST	1849152
	_		
Sequence 9165	AA232784	EST	1855986
Sequence 9166	AA233063	EST	1856075
Sequence 9167	AA233297	EST	18562 <b>9</b> 0
Sequence 9168	AA234499	EST	1858990
Sequence 9169	AA234963	EST	1858132
Sequence 9170	AA236353	EST	1858403
	AA236418	EST	
Sequence 9171			1858543
Sequence 9172	AA243646	EST	1874456
Sequence 9173	AA243672	EST	1874655
Sequence 9174	AA247773	EST	1879878
Sequence 9175	AA252002	EST	1886963
Sequence 9176	AA252445	EST	1887416
Sequence 9177	AA252465	EST	1887436
Sequence 9178	AA252567		1887541
		EST	
Sequence 9179	AA252657	EST	1887622
Sequence 9180	AA255982	EST	1891658
Sequence 9181	AA257153	EST	1891281
Sequence 9182	AA258425	EST	1893549
Sequence 9183	AA258503	EST	1893628
Sequence 9184	AA259061		1894541
		EST	
Sequence 9185	AA262003	EST	1897984
Sequence 9186	AA262559	EST	1898011
Secuence 9187	AA278264	EST	1920222
Secuence 9188	AA278642	EST	1919962
Sequence 9189	AA278910	EST	1920376
Secuence 9190	AA279735	EST	1921200
224201.00 0100		201	132 1200

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Sequence 9191	AA280710	EST	1923603
Sequence 9192	AA281238	EST	1923919
Sequence 9193	AA282006	EST	1924848
Sequence 9194	AA282870	EST	1925786
Sequence 9195	AA282920	EST	1925896
Sequence 9196	AA284386	EST	1928720
•	AA284870		
Sequence 9197		EST	1927464
Sequence 9198	AA287235	EST	1934261
Sequence 9199	AA287990	EST	1933814
Sequence 9200	AA291446	EST	1939425
Sequence 9201	AA291926	EST	1940090
Sequence 9202	AA293348	EST	1940745
•			
Sequence 9203	AA294883	EST	1947228
Sequence 9204	AA296484	₽°T	194 <b>881</b> 8
Sequence 9205	AA297767	E⊍T	19 <b>501</b> 70
Sequence 9206	AA299346	EST	1951678
Sequence 9207	AA300574	EST	1953135
Sequence 9208	AA300728	EST	1953296
Sequence 9209	AA301138	EST	1953491
Sequence 9210	AA301439	EST	1953773
Sequence 9211	AA302601	EST	1955083
Sequence 9212	AA303657	EST	1956009
Sequence 9213	AA304899	EST	1957247
Sequence 9214	AA304904	EST	1957469
Sequence 9215	AA304981	EST	1957328
Sequence 9216	AA305193	EST	1957520
•			
Sequence 9217	AA305525	EST	1957850
Sequence 9218	AA305568	EST	1957915
Sequence 9219	AA305862	EST	1958192
Sequence 9220	AA305873	EST	1958203
Sequence 9221	AA306226	EST	1958554
Sequence 9222	AA306444	EST	1958772
Sequence 9223	AA306511	EST	1958840
Sequence 9224	AA306594	EST	1958923
•			
Sequence 9225	AA306623	EST	1958952
Sequence 9226	AA306709	EST	1959037
Sequence 9227	AA306734	EST	1959083
Sequence 9228	AA306846	EST	1959174
Sequence 9229	AA306917	EST	1959247
Sequence 9230	AA306961	EST	1959291
Sequence 9231	AA306966	EST	1959316
Sequence 9232	AA307060	EST	1959538
Sequence 9233	AA307269	EST	1959597
Sequence 9234	AA307327	EST	195 <b>9655</b>
Sequence 9235	AA307332	EST	1959660
Sequence 9236	AA307779	EST	1960177
Sequence 9237	AA307876	EST	1960276
Sequence 9238	AA307979	EST	1960308
Sequence 9239	AA308390	EST	1960719
•			1960922
Sequence 9240	AA308574	EST	
Sequence 9241	AA309051	EST	1961377
Sequence 9242	AA309832	EST	1962325
Sequence 9243	AA310059	EST	1962388
Sequence 9244	AA310330	EST	1962679
Sequence 9245	AA311461	EST	1963787
Sequence 9246	AA311501	EST	1964046
22420100 0440		201	, 504040

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Sequence	9247	AA311933	EST	1964261
Sequence	9248	AA312489	EST	19 <b>6481</b> 8
Sequence		AA312669	EST	1965018
Sequence		AA312778	EST	1965126
Sequence		AA312885	EST	1965276
		AA313319		
Sequence			EST	1965648
Sequence		AA313489	EST	19 <b>6581</b> 8
Sequence		AA313896	EST	19 <b>6622</b> 5
Sequence	9255	AA314146	EST	1966495
Sequence	9256	AA314350	EST	1966679
Sequence	9257	AA314550	EST	1966899
Sequence	9258	AA314707	EST	1967056
Sequence	9259	AA314717	EST	1967277
Sequence		AA314882	EST	1967452
Sequence		AA315444	EST	1967923
Sequence		AA315472	EST	1967801
		AA315499		
Sequence			EST	1967828
Sequence		AA315975	EST	1968304
Sequence		AA316030	EST	1968359
Sequence		AA316470	EST	1968799
Sequence		AA316551	EST	1968879
Sequence		AA316590	EST	19 <b>689</b> 18
Sequence	9269	AA316620	EST	1968948
Sequence	9270	AA316641	EST	1968989
Sequence		AA316743	EST	1969142
Sequence		AA316864	EST	1969192
Sequence		AA317246	EST	1969573
Sequence		AA318236	EST	1970564
Sequence Sequence		AA318266		1970594
•			EST	
Sequence		AA323846	EST	1976173
Sequence		AA325370	EST	1977838
Sequence		AA325474	EST	1977717
Sequence:		AA326465	EST	1978709
Sequence		AA326466	EST	1978710
Sequence		AA328435	EST	1980679
Sequence	9282	AA329233	EST	1981477
Sequence:	9283	AA330793	EST	1983035
Sequence	9284	AA332288	EST	1984532
Sequence	9285	AA333116	EST	1985360
Sequence:	9286	AA335696	EST	1987937
Sequence		AA335862	EST	1988104
Sequence		AA336878	EST	1989117
Sequence		AA337404	EST	1989869
Sequence		AA337798	EST	1990057
•		AA338634		
Sequence			EST	1990882
Sequence		AA339622	EST	1991880
Sequence !		AA341987	EST	1994222
Sequence :		AA343749	EST	1995986
Sequence		AA349886	EST	2002205
Sequence !	9296	AA349983	EST	2002303
Sequence :	9297	AA351222	EST	2003561
Sequence :	9298	AA352688	EST	2005008
Sequence		AA353179	EST	2005499
Sequence		AA353913	EST	2006231
Sequence :		AA355954	EST	2008272
Sequence !		AA355956	EST	2008274
C C q u o l Co	0002	, ~~~~	E31	2000214

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Sequence 9303	AA356182	EST	2008511
			2008524
Sequence 9304	AA356195	EST	
Sequence 9305	AA358978	EST	2011296
Sequence 9306	AA359611	EST	2012169
Sequence 9307	AA359735	EST	2012052
	AA364033	EST	2016351
Sequence 9308			
Sequence 9309	AA367068	EST	2019406
Sequence 9310	AA368087	EST	2020446
Sequence 9311	AA369721	EST	2022039
Sequence 9312	AA370428	EST	2022933
Sequence 9313	AA371420	EST	2023780
		EST	2024275
Sequence 9314	AA371957		
Sequence 9315	AA372674	EST	2025182
Sequence 9316	AA375636	EST	20 <b>2795</b> 6
Sequence 9317	AA376361	EST	2028679
Sequence 9318	AA376458	EST	2028777
Sequence 9319	AA378378	EST	2030695
	AA378849	EST	2031168
Sequence 9320			2038701
Sequence 9321	AA386348	EST	
Sequence 9322	AA397884	EST	2050623
Sequence 9323	AA399142	EST	2052896
Sequence 9324	AA399177	EST	2052913
Sequence 9325	AA401252	EST	2055141
Sequence 9326	AA401261	EST	2055168
Common 0327	AA401526	EST	2053804
Sequence 9327		EST	2056854
Sequence 9328	AA402063		
Sequence 9329	AA403047	EST	2055609
Sequence 9330	AA406425	EST	2064410
Sequence 9331	AA412426	EST	2071129
Sequence 9332	AA417300	EST	2077408
Sequence 9333	AA417999	EST	2079818
Sequence 9334	AA418045	EST	2079864
	AA418096	EST	2079897
Sequence 9335			
Sequence 9336	AA418249	EST	2080078
Sequence 9337	AA421643	EST	2100459
Sequence 9338	AA424317	EST	2103287
Sequence 9339	AA425387	EST	2106170
Sequence 9340	AA425619	EST	2106375
Sequence 9341	AA425658	EST	2106433
Sequence 9342	AA425751	EST	2106453
	AA425761	EST	2106481
Sequence 9343		EST	2112054
Sequence 9344	AA427920		
Sequence 9345	AA428948	EST	2110491
Sequence 9346	AA429871	EST	2113108
Sequence 9347	AA431577	EST	2115285
Sequence 9348	AA433865	EST	2138779
Sequence 9349	AA436966	EST	2141880
	AA442297	EST	2154175
Sequence 9350			2154565
Sequence 9351	AA442687	EST	
Sequence 9352	AA446391	EST	2159056
Sequence 9353	AA448292	EST	2161962
Sequence 9354	AA449483	EST	2163233
Sequence 9355	AA449501	EST	2163251
Sequence 9356	AA449646	EST	2163396
Sequence 9357	AA450096	EST	2163846
Sequence 9358	AA450166	EST	2163916
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Sequence 9359	AA450276	EST	2164026
Sequence 9360	AA452716	EST	2166385
Sequence 9361	AA453565	EST	2167234
Sequence 9362	AA453899	EST	2167568
Sequence 9363	AA456454	EST	2179030
Sequence 9364	AA457365	EST	2180085
Sequence 9365	AA458613	EST	
Sequence 9366	AA459167	EST	2183520
Sequence 9367	AA461383	EST	2184074
Secuence 9368	AA463367	EST	2186503
Sequence 9369	AA465402		2188251
Sequence 9370	AA469033	EST	2191569
Secuence 9371	AA470067	EST	2195567
Secuence 9372		EST	2197376
Sequence 9373	AA476876	EST	2205087
Sequence 9374	AA478162	EST	2206796
	AA479499	EST	2208055
Sequence 9375	AA430816	EST	2210368
Sequence 9376	AA481233	EST	2210785
Sequence 9377	AA481838	EST	2209516
Sequence 9378	AA485255	EST	2214474
Sequence 9379	AA485520	EST	2214739
Sequence 9380	AA488447	EST	2215878
Sequence 9381	AA488488	EST	2215919
Sequence 9382	AA488513	EST	2215944
Sequence 9383	AA489117	EST	2218719
Sequence 9384	AA489288	EST	2218890
Sequence 9385	AA490389	EST	2219562
Sequence 9386	AA491213	EST	2220386
Sequence 9387	AA491638	EST	2221200
Sequence 9388	AA494334	EST	2224121
Sequence 9389	AA494355	EST	2224142
Sequence 9390	AA494387	EST	2224174
Sequence 9391	AA495912	EST	2229233
Sequence 9392 Sequence 9393	AA496486	EST	2229807
	AA501442	EST	2236409
Sequence 9394 Sequence 9395	AA504269	EST	2240429
	AA505816	EST	2241953
Sequence 9396 Sequence 9397	AA507818	EST	2244257
Sequence 9398	AA508038	EST	2244477
Sequence 9399	AA514302 AA516531	EST	2253810
Sequence 9400		EST	2256055
Sequence 9401	AA522888	EST	2263600
Sequence 9402	AA527374	EST	2269443
Sequence 9402 Sequence 9403	AA532530	EST	2276784
Sequence 9404	AA532673	EST	2276927
Sequence 9404	AA533360	EST	2277456
Sequence 9405	AA534734	EST	2278987
Sequence 9406 Sequence 9407	AA535663	EST	2279916
	AA535845	EST	2280098
Sequence 9408	AA552507	EST	2322761
Sequence 9409	AA558357	EST	2328834
Sequence 9410	AA565548	EST	2337187
Sequence 9411	AA565867	EST	2337506
Sequence 9412	AA568785	EST	2341839
Sequence 9413	AA573427	EST	2347955
Sequence 9414	AA574223	EST	2348738

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Sequence 9415	AA576016	EST	2350531
Sequence 9416	AA576889	EST	2354363
Sequence 9417	AA579611	EST	2357795
Sequence 9418	AA581185	EST	2358957
Sequence 9419	AA587264		
		EST	2398078
Sequence 9420	AA587570	EST	2397245
Sequence 9421	AA594562	EST	2409912
Sequence 9422	AA598697	EST	2432369
Sequence 9423	AA600084	EST	2433709
Sequence 9424	AA600355	EST	2433980
Sequence 9425	AA601289	EST	2434914
Sequence 9426	AA603876	EST	2437737
Sequence 9427	AA604860	EST	2445724
Sequence 9428	AA608613	EST	2457041
Sequence 9429	AA609417	EST	2457845
Sequence 9430	AA612668	EST	2463706
Sequence 9431	AA625213	EST	2537598
Sequence 9432	AA626629	EST	2539016
Sequence 9433	AA627320	EST	2540337
Sequence 9434	AA628491	EST	2540878
Sequence 9435	AA628520	EST	2540907
Sequence 9436	AA631993	EST	2555407
Sequence 9437	AA634958	EST	2558172
Sequence 9438	AA640716	EST	2565966
Sequence 9439	AA641939	EST	2567157
•	AA652388	EST	2584040
Sequence 9440			
Sequence 9441	AA659693	EST	2595847
Sequence 9442	AA662024	EST	2616115
Sequence 9443	AA663649	EST	2617640
Sequence 9444	AA668886	EST	263 <b>0385</b>
Sequence 9445	AA679400	EST	265 <b>9922</b>
Sequence 9446	AA679621	EST	2660143
Sequence 9447	AA687221	EST	2675412
Sequence 9448	AA687495	EST	2675686
Sequence 9449	AA693932	EST	2694870
Sequence 9450	AA694188	EST	2695126
Sequence 9451	AA700208	EST	2703171
Sequence 9452	AA701138	EST	2704303
Sequence 9453	AA704533	EST	2714451
Sequence 9454	AA705937	EST	2715855
Sequence 9455	AA709002	EST	2718920
Sequence 9456	AA716423	EST	2728697
Sequence 9457	AA722455	EST	2740162
Sequence 9458	AA722786	EST	2740493
Sequence 9459	AA730400	EST	2751604
Sequence 9460	AA731394	EST	275 <b>35</b> 50
Sequence 9461	AA742260	EST	2784260
Sequence 9462	AA746310	EST	2786296
Sequence 9463	AA746748	EST	2786734
Sequence 9464	AA747197	EST	2787155
Sequence 9465	AA749387	EST	278 <b>93</b> 45
Sequence 9466	AA757433	EST	280 <b>529</b> 6
Sequence 9467	AA761101	EST	28 <b>1003</b> 1
Sequence 9468	AA765907	EST	2817145
Sequence 9469	AA769588	EST	2820826
Sequence 9470	AA772570	EST	28 <b>24353</b>

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Sequence 9471	AA774779	EST	2834113
Secuence 9472	AA775625	EST	2834959
Secuence 9473	AA805474	EST	2874224
Sequence 9474	AA808205	EST	2877611
Sequence 9475	AA808239	EST	2877645
Sequence 9476	AA813991	EST	2883587
Sequence 9477	AA814182	EST	2883778
Sequence 9478	AA814817	EST	2884413
Sequence 9479	AA826906	EST	2900903
Sequence 9480	AA830072	EST	2903171
Sequence 9481	AA831991	EST	2905090
Sequence 9482	AA833749	EST	2908517
Sequence 9483	AA837407	EST	2912606
Sequence 9484	AA838157	EST	2913814
Sequence 9485	AA843207	EST	2929725
Sequence 9486	AA843238	EST	2929756
Sequence 9487	AA843878	EST	2930329
Sequence 9488	AA843908	EST	2930359
Sequence 9489	AA857964	EST	29 <b>4626</b> 6
Sequence 9490	AA864964	EST	29 <b>5927</b> 7
Sequence 9491	AA868452	EST	29 <b>6389</b> 7 -
Sequence 9492	AA868722	EST	2964167
Sequence 9493	AA872507	EST	2968685
Sequence 9494	AA872913	EST	2969035
Sequence 9495	AA872991	EST	2969113
Sequence 9496	AA877350	EST	29 <b>8642</b> 7
Sequence 9497	AA877789	EST	29 <b>86754</b>
Sequence 9498	AA902103	EST	3037293
Sequence 9499	AA902974	EST	3038097
Sequence 9500	AA903253	EST	3038376
Sequence 9501 Sequence 9502	AA905330	EST	3040453
Sequence 9503	AA906868 AA907186	EST	3042112
Sequence 9504	AA908460	EST	3042646
Sequence 9505	AA913203	EST	3047865
Sequence 9506	AA916003	EST EST	3052595
Sequence 9507	AA927612	EST	305 <b>539</b> 5 30 <b>77033</b>
Sequence 9508	AA931566	EST	308 <b>5</b> 9 <b>5</b> 2
Sequence 9509	AA932278	EST	3086516
Sequence 9510	AA968524	EST	3143704
Sequence 9511	AA973790	EST	3148970
Sequence 9512	AA976114	EST	3151906
Sequence 9513	AA976653	EST	3154099
Sequence 9514	AA984122	EST	3162647
Sequence 9515	AA988106	EST	3173470
Sequence 9516	AA988525	EST	3174217
Sequence 9517	AA988576	EST	3174268
Sequence 9518	AA991580	EST	3178069
Sequence 9519	AF001542	EST	2529714
Sequence 9520	AI015791	EST	3230127
Sequence 9521	AI017095	EST	3231431
Sequence 9522	Al017902	EST	3232238
Sequence 9523	Al018625	EST	3233144
Sequence 9524	AI022209	EST	3239562
Sequence 9525	AI027035	EST	3244551
Sequence 9526	AI033991	EST	3254944

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Sequence 9527	AI034371	EST	3255324
Sequence 9528	AI040602	EST	3279796
Sequence 9529	AI051466	EST	3307000
Sequence 9530	AI052000	ËST	3307991
			3330203
Sequence 9531	AI056337	EST	
Sequence 9532	AI056402	EST	3330268
Sequence 9533	Al061175	EST	3336543
Sequence 9534	AI064910	EST	6359182
Sequence 9535	AI066583	EST	3367285
Sequence 9536	AI076630	EST	3405808
Sequence 9537	AI076953	EST	3404782
Sequence 9538	AI078217	EST	3412625
Sequence 9539	AI079510	EST	3415761
Sequence 9540	Al080497	EST	3416748
Sequence 3340	Al081780	EST	3418572
Sequence 9541	AI081780	EST	3423217
Sequence 9542			
Sequence 9543	AI084916	EST	3423339
Sequence 9544	Al085616	EST	3424039
Sequence 9545	AI088897	EST	3427956
Sequence 9546	AI090197	EST	3429256
Sequence 9547	Al091173	EST	3430232
Sequence 9548	Al094514	EST	3433490
Sequence 9549	AI095450	EST	3434426
Sequence 9550	Al114692	EST	6360037
Sequence 9551	Al123418	EST	3539184
Sequence 9552	Al124792	EST	3593306
Sequence 9553	Al126005	EST	3594519
Sequence 9554	Al129250	EST	3597764
Sequence 9555	Al133074	EST	6360390
	Al133088	EST	6360404
Sequence 9556	Al133274	EST	6360590
Sequence 9557			6360742
Sequence 9558	Al133426	EST	3645736
Sequence 9559	AI139764	EST	
Sequence 9560	Al142414	EST	3658773
Sequence 9561	AI146833	EST	3674515
Sequence 9562	Al148106	EST	3675788
Sequence 9563	AI150430	EST	3678899
Sequence 9564	Al151146	EST	3679615
Sequence 9565	Al159774	EST	3693133
Sequence 9566	Al161072	EST	3694377
Sequence 9567	AI168316	EST	3701486
Sequence 9568	AI168348	EST	3701518
Sequence 9569	AI186453	EST	3737091
Sequence 9570	AI187157	EST	3737795
Sequence 9571	AI189475	EST	3740684
Sequence 9572	AI193372	EST	3744581
Sequence 9573	Al198218	EST	3750824
Sequence 9574	Al199191	EST	3751797
Sequence 9575	Al199194	EST	3751800
	Al202137	EST	3754743
Sequence 9576			3755820
Sequence 9577	Al203214	EST	
Sequence 9578	AI207471	EST	6361479
Sequence 9579	Al207868	EST	3769810
Sequence 9580	AI214048	EST	3777649
Sequence 9581	Al214453	EST	3778054
Sequence 9582	Al216098	EST	3785139

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Sequence !	9583	AI217246	EST	3797061
Sequence		AI244268	EST	3839665
Sequence !		Al248386	EST	3843783
Sequence !		A1249242	EST	3845771
Sequence :	9587	AI251739	EST	3848268
Sequence !		AJ261980	EST	3870183
Sequence !		Al262251	EST	3870454
		AI267162	EST	
Sequence 9				3886329
Sequence !		Al267848	EST	3887015
Sequence !		Al268335	EST	3887502
Sequence !	9593	Al268362	EST	3887529
Sequence !	9594	AI270685	EST	3889852
Sequence 9		Al275466	EST	3897740
Sequence !		Al276933	EST	3899201
Sequence 9		AI277443	EST	3899711
Sequence		AI278328	EST	3900596
Sequence !		Al279625	EST	3917859
Sequence !	9600	AI280561	EST	3918794
Sequence !	9601	Al281394	EST	3919627
Sequence !		Al283373	EST	3921606
Sequence 9		AI283578	EST	3921811
Sequence !		AI284482	EST	
				3922715
Sequence 9		Al290052	EST	3931718
Sequence !		Al290111	EST	3931777
Sequence !	9607	Al290154	EST	3931820
Sequence !	9608	Al291087	EST	3933861
Sequence !	9609	Al292094	EST	3934868
Sequence 9	9610	Al298974	EST	3958628
Sequence 9		Al333116	EST	4069675
Sequence		Al334177	EST	4070736
		Al334889	EST	
Sequence !				4071816
Sequence !		Al337358	EST	4074285
Sequence !		Al339284	EST	4076211
Sequence 9	9616	Al341138	EST	4078065
Sequence !	9617	Al341291	EST	4078218
Sequence 9	9618	Al342090	EST	4079017
Sequence !	9619	Al343404	EST	4080610
Sequence !		Al344189	EST	4081395
Sequence !		Al348588	EST	4085794
Sequence !		Al349337	EST	4086543
Sequence !		Al349957	EST	4087163
Sequence !		Al350524	EST	4087730
Sequence 9	9625	Al352184	EST	4089390
Sequence !	9626	Al355277	EST	4095430
Sequence!	9627	Al356461	EST	4108082
Sequence !		Al357566	EST	4109187
Sequence !		Al357582	EST	4109203
Sequence !		Al360681		4112302
			EST	
Sequence 9		Al361796	EST	4113417
Sequence !		Al373828	EST	4153694
Sequence !		Al378900	EST	4188753
Sequence !	9634	Al379649	EST	4189502
Sequence !		Al381737	EST	4194518
Sequence !		Al381972	EST	4194753
Sequence 9		Al393269	EST	4222816
		Al420586	EST	4266517
Sequence 9	5000	M420000	L31	4200017

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Sequence 9639	Al435112	EST	4300898
Sequence 9640	Al439421	EST	4304204
Sequence 9641	Al439588	EST	4305374
Sequence 9642	Al445716	EST	4290157
Sequence 9643	Al453049	EST	4307708
Companie 9043	Al457432	EST	4310301
Sequence 9644			4310833
Sequence 9645	AI458254	EST	4311178
Sequence 9646	Al458599	EST	
Sequence 9647	Al458758	EST	4311337
Sequence 9648	Al460220	EST	4313101
Sequence 9649	Al468419	EST	4330509
Sequence 9650	A1472084	EST	4334174
Sequence 9651	Al472566	EST	4325611
Sequence 9652	Al479075	EST	4372243
Sequence 9653	A1480254	EST	4373422
Sequence 9654	Al492032	EST	4393035
Sequence 9655	Al494037	EST	4395040
Sequence 9656	AI525541	EST	4439676
Sequence 9657	AI525654	EST	4439789
Sequence 9658	AI538842	EST	4452977
Sequence 9659	AI557226	EST	4489589
Sequence 9660	AI560973	EST	4511314
Sequence 9661	AI563921	EST	4522378
Sequence 9662	Al564719	EST	4523176
Sequence 9002	AI570615	EST	4533989
Sequence 9663	AI580252	EST	4564628
Sequence 9664	AI580336	EST	4564712
Sequence 9665	Al581098	EST	4565474
Sequence 9666	Al597918	EST	4606977
Sequence 9667		EST	4617963
Sequence 9668	Al608796	EST	4618254
Sequence 9669	Al609087	EST	4620047
Sequence 9670	A1610880		4648915
Sequence 9671	AI623984	EST EST	4649682
Sequence 9672	AI624751		4665154
Sequence 9673	Al628354	EST	4665217
Sequence 9674	Al628417	EST	4683800
Sequence 9675	Al632470	EST	
Sequence 9676	Al633000	EST	4684330
Sequence 9677	AI637826	EST	4690060
Sequence 9678	Al638561	EST	4690795
Sequence 9679	AI640607	EST	4703716
Sequence 9680	Al650583	EST	4734562
Sequence 9681	AI651877	EST	4735856
Sequence 9682	Al672492	EST	4852223
Sequence 9683	A1684595	EST	4895889
Sequence 9684	A1685930	EST	4897224
Sequence 9685	A1690089	EST	4901383
Sequence 9686	A1692658	EST	4969998
Sequence 9687	Al694944	EST	4982844
Sequence 9688	Al697185	EST	4985085
Sequence 9689	Al720354	EST	5037610
Sequence 9690	A1720743	EST	5037999
Sequence 9691	AI738670	EST	5100651
Sequence 9692	AI743182	EST	5111470
Sequence 9693	AI743921	EST	5112209
Sequence 9694	AI750105	EST	5128369

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Sequence 9695	AI751208	EST	5129472
Sequence 9696	AI751459	EST	5129723
Sequence 9697	A1751487	EST	5129751
Sequence 9698	A1752264	EST	5130528
Sequence 9699	A1752414	EST	5130678
Sequence 9700	AI752425	EST	5130689
Sequence 9701	AI760045	EST	5175712
Sequence 9702	AI761315	EST	5176982
Sequence 9703	AI765501	EST	5232010
Sequence 9704	AI766305	EST	
			5232814
Sequence 9705	AI767578	EST	5234087
Sequence 9706	AI767726	EST	5234235
Sequence 9707	AI792015	EST	5339731
Sequence 9708	AI795827	EST	5361287
Sequence 9709	Al797228	EST	5362700
Sequence 9710	AI800215	EST	5365687
Sequence 9711	AI806017	EST	5392583
Sequence 9712	AI807798	EST	5394364
Sequence 9713	AI808865	EST	5395431
Sequence 9714	Al810669	EST	5397235
Sequence 9715	Al814463	EST	5425678
Sequence 9716	Al814513	EST	5425728
Sequence 9717	Al815492	EST	5431038
Sequence 9718	AI816438	EST	5431984
Sequence 9719	AI818683	EST	5437762
Sequence 9720	Al819405	EST	5438484
Sequence 9721	AI828075	EST	5448746
Sequence 9722	AI863382	EST	5527489
Sequence 9723	AI873001	EST	5547050
Sequence 9724	AI873501	EST	5547550
Sequence 9725	A1874326	EST	5548375
Sequence 9726	AI884491	EST	5589655
Sequence 9727	A1884783	EST	5589947
Sequence 9728	AI885021	EST	5590185
Sequence 9729	AI887758	EST	5592922
Sequence 9730	AI888327	EST	5593577
Sequence 9731	AI888872	EST	5594036
Sequence 9732	AI903705	EST	6494092
Sequence 9733	AI904447	EST	6494834
	AI906328		
Sequence 9734		EST	6496715
Sequence 9735	AI910090	EST	6500770
Sequence 9736	AI914133	EST	5633988
Sequence 9737	Al918605	EST	5638460
Sequence 9738	Al924534	EST	5660498
Sequence 9739	AI927755	EST	5663719
Sequence 9740	AI929696	EST	5665660
Sequence 9741	AI936166	EST	5675036
Sequence 9742	AI950991	EST	5743301
Sequence 9743	AI970095	EST	5766921
Sequence 9744	Al972423	EST	5769339
Sequence 9745	AI982834	EST	5810053
Sequence 9746	Al982890	EST	5810109
Sequence 9747	AL035972	EST	5405602
Sequence 9748	AL036220	EST	5405817
Sequence 9749	AL036415	EST	5405971
Sequence 9750	AL037386	EST	5406786
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Sequence 9751	AL037968	EST	5928298
Sequence 9752	AL039569	EST	5408608
Sequence 9753	AL039961	EST	5408942
Sequence 9754	AL040667	EST	5409613
Sequence 9755	AL040692	EST	5409638
Sequence 9756	AL041673	EST	5421022
Sequence 9757	AL042613	EST	5935505
			5422129
Sequence 9758	AL042680	EST	
Sequence 9759	AL042725	EST	5422174
Sequence 9760	AL042986	EST	59 <b>356</b> 55
Sequence 9761	AL043210	EST	5935803
Sequence 9762	AL043542	EST	5422929
Sequence 9763	AL043672	EST	5423059
0	AL045757	EST	5936079
Sequence 9764			
Sequence 9765	AL045793	EST	5433905
Sequence 9766	AL045936	EST	5434039
Sequence 9767	AL045937	EST	5936123
Sequence 9768	AL045976	EST	5 <b>4340</b> 76
Sequence 9769	AL046015	EST	5434109
	AL047001	EST	5435057
Sequence 9770			4727279
Sequence 9771	AL047332	EST	
Sequence 9772	AL047890	EST	4728078
Sequence 9773	AL048554	EST	5936 <b>5</b> 65
Sequence 9774	AL110421	EST	5 <b>86602</b> 9
Sequence 9775	AL118756	EST	59 <b>2465</b> 5
Sequence 9776	AL118845	EST	5924744
Sequence 9777	AL119809	EST	5925708
		EST	5926610
Sequence 9778	AL120711		
Sequence 9779	AL120865	EST	5926866
Sequence 9780	AL120911	EST	5926912
Sequence 9781	AL121473	EST	592 <b>7474</b>
Sequence 9782	AL121558	EST	5927559
Sequence 9783	AL133779	EST	6601967
Sequence 9784	AL133900	EST	6602087
Sequence 9785	AL134724	EST	6602911
		EST	6602982
Sequence 9786	AL134795		
Sequence 9787	AL134938	EST	6603125
Sequence 9788	AL135007	EST	6603194
Sequence 9789	AL135023	EST	6603210
Sequence 9790	AL135290	EST	6603477
Sequence 9791	AL135343	EST	6603530
Sequence 9792	AL135362	EST	6603549
Sequence 9793	AL135502	EST	6603689
Sequence 9793			6603787
Sequence 9794	AL135600	EST	
Sequence 9795	AL138042	EST	6854722
Sequence 9796	AW002442	EST	5849358
Sequence 9797	AW014499	EST	5863256
Sequence 9798	AW014727	EST	58 <b>63484</b>
Sequence 9799	AW029611	EST	5888367
Sequence 9800	AW029642	EST	5888398
Coquente 3000	AW043897	EST	5904426
Sequence 9801			6014222
Sequence 9802	AW062837	EST	· ·
Sequence 9803	AW068104	EST	6023102
Sequence 9804	AW070431	EST	602 <b>5429</b>
Sequence 9805	AW070690	EST	6025688
Sequence 9806	AW080672	EST	6035905
20420.00			

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Sequence 9807	AW084080	EST	603 <b>923</b> 2
Sequence 9808	AW118447	EST	5087031
Sequence 9809	AW148635	EST	6196531
Sequence 9810	AW149925	EST	6197 <b>821</b>
Sequence 9811	AW157231	EST	622 <b>863</b> 2
Sequence 9812	AW160468	EST	629 <b>95</b> 01
Sequence 9813	AW160833	EST	629 <b>986</b> 6
Sequence 9814	AW161807	EST	630 <b>084</b> 0
Sequence 9815	AW161978	EST	6301 <b>011</b>
Sequence 9816	AW162587	EST	6301 <b>62</b> 0
Sequence 9817	AW162761	EST	6301794
Sequence 9818	AW162774	EST	6301807
Sequence 9819	AW163485	EST	6302518
Sequence 9820	AW163783	EST	6302816
Sequence 9821	AW175627	EST	6441768
Sequence 9822	AW177931	EST	6443968
	AW178072	EST	6444109
Sequence 9823			
Sequence 9824	AW194013	EST	6472742
Sequence 9825	AW237543	EST	656 <b>9932</b>
Sequence 9826	AW239041	EST	6571431
Sequence 9827	AW239281	EST	6571 <b>6</b> 71
Sequence 9828	AW245631	EST	658 <b>8624</b>
Sequence 9829	AW245889	EST	658 <b>8882</b>
Sequence 9830	AW245906	EST	658 <b>8899</b>
Sequence 9831	AW245923	EST	658 <b>8916</b>
Sequence 9832	AW246049	EST	6589042
Sequence 9833	AW246074	EST	65 <b>89067</b>
Sequence 9834	AW247459	EST	6590452
Sequence 9835	AW247494	EST	6590487
Sequence 9836	AW249281	EST	6592274
Sequence 9837	AW250125	EST	6593118
Sequence 9838	AW250126	EST	6593119
	AW263239	EST	6640055
Sequence 9839			
Sequence 9840	AW270779	EST	6657809
Sequence 9841	AW273055	EST	6660085
Sequence 9842	AW274583	EST	6661613
Sequence 9843	AW292896	EST	6699532
Sequence 9844	AW292897	EST	669 <b>9533</b>
Sequence 9845	AW293159	EST	6699795
Sequence 9846	AW294649	EST	6701285
Sequence 9847	AW294971	EST	6701 <b>607</b>
Sequence 9848	AW296629	EST	6703 <b>26</b> 5
Sequence 9849	AW303456	EST	6713145
Sequence 9850	AW327374	EST	6797869
Sequence 9851	AW327483	EST	6797 <b>978</b>
Sequence 9852	AW340172	EST	6836798
Sequence 9853	AW340708	EST	6837334
Sequence 9854	AW360965	EST	6865615
Sequence 9855	AW361001	EST	686 <b>565</b> 1
Sequence 9856	AW361103	EST	6865753
Sequence 9857	AW361711	EST	6866361
Sequence 9858	AW362982	EST	6867632
Sequence 9859	AW363561		6868211
		EST	
Sequence 9860	AW364748	EST	6869502
Sequence 9861	AW364827	EST	68 <b>69386</b>
Sequence 9862	AW365085	EST	68 <b>69735</b>

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Sequence 9863	AW365852	EST	6870502
Sequence 9864	AW368293	EST	6872943
Sequence 9865	AW368461	EST	6873111
Sequence 3003	AW375589	EST	6880152
Sequence 9866		EST	6881580
Sequence 9867	AW376917	EST	6884486
Sequence 9868	AW379827		6885886
Sequence 9869	AW381227	EST	
Sequence 9870	AW381248	EST	6885907
Sequence 9871	AW383599	EST	6888167
Sequence 9872	AW384101	EST	6888760
Sequence 9873	AW384915	EST	6889574
Sequence 9874	AW385192	EST	6889851
Sequence 9875	AW385322	EST	6889981
Sequence 9876	AW385328	EST	6889987
Sequence 9877	AW388939	EST	6893598
Sequence 9878	AW392421	EST	6897080
Sequence 9879	AW393095	EST	6897754
Sequence 9079	AW401915	EST	6920601
Sequence 9880	AW402136	EST	6920822
Sequence 9881	AW402676	EST	6921388
Sequence 9882	AW404132	EST	6923189
Sequence 9883	AW405567	EST	6924624
Sequence 9884		EST	6926269
Sequence 9885	AW407212	EST	6927331
Sequence 9886	AW408274		6927480
Sequence 9887	AW408423	EST	6936061
Sequence 9888	AW410520	EST	
Sequence 9889	AW410772	EST	6936313 6936752
Sequence 9890	AW411211	EST	
Sequence 9891	AW411217	EST	6936758
Sequence 9892	AW474433	EST	7044539
Sequence 9893	AW498895	EST	7137236
Sequence 9894	AW500117	EST	7112425
Sequence 9895	AW500758	EST	7113663
Sequence 9896	AW501342	EST	7114784
Sequence 9897	AW501373	EST	7114844
Sequence 9898	AW501390	EST	7114878
Sequence 9899	AW501401	EST	7114900
Sequence 9900	AW501421	EST	7114936
Sequence 9901	AW515416	EST	N/A
Sequence 9902	AW578462	EST	7253511
Sequence 9903	AW578582	EST	7253631
Sequence 9904	AW582604	EST	7257653
Sequence 9905	AW583629	EST	7260587
Sequence 9906	C04057	EST	1467308
Sequence 9907	F06849	EST	672465
Sequence 9908	F09531	EST	682064
Sequence 9909	F11462	EST	705760
Sequence 9910	F12434	EST	706790
Sequence 9911	H01873	EST	864806
Sequence 9912	H08043	EST	872865
Sequence 9913		EST	872915
Sequence 9914		EST	874050
Sequence 9915		EST	874139
Sequence 9916		EST	874615
Sequence 9917		EST	875383
Sequence 9918		EST	875713
Sedneuce 2210	,,,,,,,,,		

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Sequence 9919	H11277	EST	876097
Sequence 9920	H15211	EST	880031
Sequence 9921	H168 <b>65</b>	EST	883105
Sequence 9922	H22726	EST	891421
Sequence 9923	H26047	EST	895170
Sequence 9924	H299 <b>98</b>	EST	900908
Sequence 9925	H30612	EST	901522
Sequence 9926	H62143	EST	1014975
Sequence 9927	H69585	EST	1039791
Sequence 9928	H70297	EST	1040503
Sequence 9929	H86644	EST	1068223
Sequence 9930	H99096	EST	1123764
Sequence 9931	M85706	EST	274353
Sequence 9932	N26971	EST	1141319
Sequence 9933	N28905	EST	1147141
Sequence 9934	N28944	EST	1147180
Sequence 9935	N34003	EST	1154403
Sequence 9936	N34221	EST	1155363
Sequence 9937	N36764	EST	1157906
Sequence 9938	N42722	EST	1167152
Sequence 9939	N43012	EST	1166756
Sequence 9940	N46795	EST	1187961
Sequence 9941	N54326	EST	1195646
Sequence 9942	N57253	EST	1201143
Sequence 9943	N95531	EST	1267800
Sequence 9944	R17855	EST	771465
Sequence 9945	R19920	EST	774554
Sequence 9946	R34894	EST	79 <b>179</b> 5
Sequence 9947	R35355	EST	792256
Sequence 9948	R35487	EST	792388
Sequence 9949	R50695	EST	812597
Sequence 9950	R52763	EST	814665
Sequence 9951	R53255	EST	815157
Sequence 9952	R59084	EST	829779
Sequence 9953	R69898	EST	843415
Sequence 9954	R73759	EST	848129
Sequence 9955	R84545	EST	942951
Sequence 9956	R88895	EST	953722
Sequence 9957	R89026	EST	953853
Sequence 9958 Sequence 9959	R93920	EST	969315
Sequence 9960	T05548	EST	316698
Sequence 9961	T08921	EST	389949
Sequence 9962	T11899 T18975	EST	596603
Sequence 9963		EST	601018
Sequence 9964	T19025	EST	601068
Sequence 9965	T32930	EST	615028
Sequence 9966	T63081	EST	666738
Sequence 9967	T77019	EST	694222
Sequence 9968	T84756	EST	713108
Sequence 9969	W02349	EST	1274356
Sequence 9970	W03691	EST	1275555
Sequence 9971	W07619	EST	1281621
Sequence 9972	W27890	EST	1307838
Sequence 9973	W28867	EST	1308878
Sequence 9974	W38913	EST	1320619
Dequence 33/4	W38915	EST	1320622

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Sequence 9975	W39482	EST	1321190
Sequence 9976	W52153	· EST	1349763
	W60983	EST	1367742
Sequence 9977			
Sequence 9978	W69356	EST	1378667
Sequence 9979	W72044	EST	1382314
Sequence 9980	W76216	EST	1386441
Sequence 9981	W79237	EST	1390368
1			
Sequence 9982	W80613	EST	1391700
Sequence 9983	W81613	EST	1392652
Sequence 9984	W88815	EST	1404287
Sequence 9985	W95032	EST	1424152
Sequence 9986	Z25224	EST	395863
		EST	56 <b>584</b> 4
Sequence 9987	Z42427		
Sequence 9988	Z45029	EST	574221
Sequence 9989	Z45988	EST	575222
Sequence 9990	Z78343	EST	1495116
Sequence 9991	Q46540	NUCPATENT	N/A
,		NUCPATENT	N/A
Sequence 9992	Q84560		
Sequence 9993	T33624	NUCPATENT	N/A
Sequence 9994	T71086	NUCPATENT	N/A
Sequence 9995	T94119	NUCPATENT	N/A
Sequence 9996	V04699	NUCPATENT	N/A
Sequence 9997	V05728	NUCPATENT	N/A
Sequence 9998	V34175	NUCPATENT	N/A
,		NUCPATENT	N/A
Sequence 9999	V34185		
Sequence 10000	V35457	NUCPATENT	N/A
Sequence 10001	V36 <b>3</b> 03	NUCPATENT	N/A
Sequence 10002	V41451	NUCPATENT	N/A
Sequence 10003	V58631	NUCPATENT	N/A
Sequence 10004	V84411	NUCPATENT	N/A
Sequence 10005	V84546	NUCPATENT	N/A
Sequence 10006	V87058	NUCPATENT	N/A
		NUCPATENT	N/A
Sequence 10007	V89685		
Sequence 10008	V90259	NUCPATENT	N/A
Sequence 10009	V99828	NUCPATENT	N/A
Sequence 10010	X00616	NUCPATENT	N/A
Sequence 10011	X05715	NUCPATENT	N/A
Sequence 10012	X37479	NUCPATENT	N/A
Sequence 10013	X40381	NUCPATENT	N/A
Sequence 10013	X61430	NUCPATENT	N/A
			N/A
Sequence 10015	X77342	NUCPATENT	
Sequence 10016	X80672	NUCPATENT	N/A
Sequence 10017	X87868	NUCPATENT	N/A
Sequence 10018	X91484	NUCPATENT	N/A
Sequence 10019	X97576	NUCPATENT	N/A
Sequence 10020	X97718	NUCPATENT	N/A
Sequence 10021	X97772	NUCPATENT	N/A
			N/A
Sequence 10022	X98309	NUCPATENT	
Sequence 10023	X98400	NUCPATENT	N/A
Sequence 10024	X98781	NUCPATENT	N/A
Sequence 10025	X99117	NUCPATENT	N/A
Sequence 10026	Z00428	NUCPATENT	N/A
Sequence 10027	Z14314	NUCPATENT	N/A
Sequence 10027	Z14685	NUCPATENT	N/A
	Z15023	NUCPATENT	N/A
Sequence 10029	Z13023 Z24893		N/A
Sequence 10030	ZZ4583	NUCPATENT	1N/PA

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# Table 3-3

Sequence 10031	Z27237	NUCPATENT	N/A
Sequence 10032	Z27237 Z29638	NUCPATENT	N/A N/A
•			
Sequence 10033	Z38853	NUCPATENT	N/A
Sequence 10034	Z39903	NUCPATENT	N/A
Sequence 10035	Z41312	NUCPATENT	N/A
Sequence 10036	Z52414	NUCPATENT	N/A
Sequence 10037	Z52416	NUCPATENT	N/A
Sequence 10038	Z53013	NUCPATENT	N/A
Sequence 10039	Z57842	NUCPATENT	N/A
Sequence 10040	Z60383	NUCPATENT	N/A
Sequence 10041	Z65271	NUCPATENT	N/A
Sequence 10042	Z80193	NUCPATENT	N/A
Sequence 10043	Z80472	NUCPATENT	N/A
Sequence 10044	Z97065	NUCPATENT	N/A
Sequence 10045	Z97145	NUCPATENT	N/A
Sequence 10046	Z97322	NUCPATENT	N/A
Sequence 10047	ACC5588	PREPATNUC	N/A
Sequence 10048	AC37772	PREPATNUC	N/A
Sequence 10049	AC38092	PREPATNUC	N/A
Sequence 10050	AC38120	PREPATNUC	N/A
Sequence 10051	AC38948	PREPATNUC	N/A
Sequence 10052	AC39251	PREPATNUC	N/A
Sequence 10053	AC39528	PREPATNUC	N/A
Sequence 10054	AC39561	PREPATNUC	N/A
,			

BNSD00 D VAC 1153836AC

## Table 3-4

Sequence 10047: Found in patent publication WO99/09061 CCCGCGGTGGCCGAGGTACTGCGTGGCCGTCGCTGTTGTGTGGCCTGGCCGCAA

Sequence 10048: Found in patent publication WO00/06698

Sequence 10049: Found in patent publication WO00/06698

TGATCTTCAGCTGCTCCTGAACTTTGGCACTGTAAGGGTTCTCTGTATCGTATGGTGGTG ACAAAGCTTTTAAAATAACCTNTCCTCCAGTGCATGGAAAAGGACTGGAGTATTTATNAA GT

Sequence 10050: Found in patent publication WO00/06730

Sequence 10051: Found in patent publication WO00/11171

Sequence 10052: Found in patent publication WO00/15799

ACCGCGGGGGCGCCGNGGNTTCNGCCTGGAGGAGCTCAGGGTGGCCGGCATTCACAAG

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# <u>Table 3-4</u>

ANAAACACTTTGCTTTGGACTAGTTAAAGCCAANAATTTGA
Sequence 10053: Found in patent publication WO00/06698
ACATCTTCCAGGACAAGGTAAACTCTGACATGCACTAGGTATGTGCAGATCCCGGCCCCT
GCCACCCAGCCTCATGCAAGTCATCCCCGACATGCACGACCGCAATGCAAGGAGG
GGAAGAAAGTCACAGCACTGATGAGGACAGCTGCAGAGGTGGCAGTGTGGACACAGGA
AGTTTGGGCCCCCTCCCTGCCCCAGCTTTCCTAGGCCAGAATTGTTTTGGCAGTAATTG
TCTGTTTAAAAAAATAAAAAGGAGAGAGGAAGCGTTCACCGCCACAAATCATAAAATGGACA
TGACTGTGGAGTCTTACAGTTCAGGGTTCTTTCATTCACGTCCCTTCCTGTCTCGGTCTG
CGGTCTTTACCACATCAATAGGACTTTTTATGCGTCCGGGTTAATTTTTCACTCCAGTGC
GTCCTGTTGCAGGGACCGGAGCTGATGGGAGCTGCTTCCCCCATGCCTCACTGGTCCCA
GATCANGGCTTCAGGGACAGATGATGAGTCTCAAACGAGCCANCAGGGGTTCTTTTGGGT
ATAAATGGGCAATTCGNCCTGTCTTAAGNCTGATGACCTCANCCGTGGTTTTTTGGAT
Sequence 10054: Found in patent publication WO00/18922
TAGGGCGAATTGGAGCTCCCCGCGGTGGCGGCCGAGGTACCTGAGCCAGGAGGAGGCCC

GGCCGTGGACCAGGAGCTATTTAACGAATACCAGTTCAGCGTGGACCAACTTATGGAACT
GGCCGGGCTGAGCTGTGCTACAGCCATCGCCAAGGCATATCCCCCCACGTCCATGTCCAG
GAGCCCCCCTACTGTCCTGGTCATCTGTGGCCCGGGGAATAATGGAGGAGATGGTCTGGT
CTGTGCTCGACACCTCAAACTCTTTGGCTACGAGCCAACCATCTATTACCCCAAAAGGCC
TAACAAGCCCTCTTCACTGCATTGGTGACCCAGTGTCAGAAAATGGACATCCCTTTCCT
TGGGGAAATGCCCGCAGAGCCCATGACGATTGATGAACTGTTGAGCTGGTGGATGCC
ATCTTTGGCTTCAAGGGCGATGTTCGGGAAC

## **Table 3-4**

Sequence 10047: Found in patent publication WO99/09061 CCCGCGGTGGCCGAGGTACTGCGTGGCCGTCCGCTGTTGTGTGGCCTGGGCCGCAA

Sequence 10048: Found in patent publication WO00/06698

CCGCGCTGCCGACGCCGACGTCTGAACTGCGGGGTCTCTATCGCACTGCTAGGGGTTCTGCT GCTGGGTGCGCCCCCCGCCGCGGGGCAGAAGCTTTTGAGATTGCTCTGCCACGAGA AAGCAACATTACAGTTCTCATAAAGCTGGGGACCCCGACTCTGCTGGCAAAACCCTGTTA CATCGTCTTTTCTAAAAGACATATAACCATGTTGTCCATCAAGTCTGGAGAAAAGAATAGT CTTTACCTTTAGCTGCCAGAGTCCTGAGAATCACTTTGTCATANAGATCCAGAAAAATAT TGACTGTATGTCAGGCCCATGTCCTTTTGGGGAGGTTCAGCTTCAGCCCTCGACATCGTT GTTGCCTACCCTCAACAGAACTTTCATCTGGGATGTCAAA

Sequence 10049: Found in patent publication WO00/06698

TGATCTTCAGCTGCTCCTGAACTTTGGCACTGTAAGGGTTCTCTGTATCGTATGGTGGTG ACAAAGCTTTTAAAATAACCTNTCCTCCAGTGCATGGAAAAGGACTGGAGTATTTATNAA GT

Sequence 10050: Found in patent publication WO00/06730

Sequence 10051: Found in patent publication WO00/11171

Sequence 10052: Found in patent publication WO00/15799

ACCGCGGGGGCCGNGGNTTCNGCCTGGAGGAGCTCAGGGTGGCCGGCATTCACAAG

A
AGGTGGCCCGGACCATCGGCATTTCTGTGGATCCGAGGAGGCGGAACAAGTCCACGGAGT
CCCTGCAGACCAACGTGCAGCGGCTGAAGGAGTACCACAAAGGGACCCAAATTCAGCGGT
CTGTGCCTACAAACTTCATTAATAACTGCTTGCAGATTGGCAGCTATCTGGTCACTTGAC
ATATCCAATGTTGCTATTTTGGTCTGGAGAAAGTTCTCCCTTTCTTCATCTACCTTAATT
TCATGTCCATTTTTAAATAATTCAAACATTTTGGGGATGTCACGGCCAATGGAATTTCGA
GAAAGCTTTGGATATTTTTTATTCAGTTTCTTCCTTAATCGATTAAGTTCAGGCATTATT
TCTGGAACAGCTACGTAAAAGTCTGCAACAATTTCATCATCCCAAATCTTCTGTATCAGA
CTAAGTGCCTCCTGCAAATGCAGCTCCATTTTCTTCCGCTATTTTGACCTCTGATGCATT
CTCTGTAAATACAGCAACTTTATTGATTTCCGAAGCCAATGGGTATGGCAAACTAANAAC
ACTGGTAAATGGCTTCACGTTTTTCTTNTTTTCCAGTGCCATATNCAGTGTCAAAATCAA

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# **Table 3-4**

ANAAAACACTTTGCTTTGGACTAGTTAAAGCCAANAATTTGA Sequence 10053: Found in patent publication WO00/06698 ACATOTTCCAGGACAAGGTAAACTCTGACATGCACTAGGTATGTGCAGATCCCGGCCCCT GCCACCCAGCCTCATGCAAGTCATCCCCGACATGACCTTCACGACCGCAATGCAAGGAGG GGAAGAAGTCACAGCACTGATGAGGACAGCTGCAGAGGTGGCAGTGTGTGGACACAGGA AGTTTGGGCCCCCTCCCTGCCCCAGCTTTCCTAGGCCAGAATTGTGTTTGGCAGTAATTG TCTGTTTAAAAAAATAAAAAGGAGGAGGAGCGTTCACCGCCACAAATCATAAAATGGACA TGACTGTGGAGTCTTACAGTTCAGGGTTCTTTCATTCACGTCCCTTCCTGTCTCGGTCTG CGGTCTTTACCACATCAATAGGACTTTTTATGCGTCCGGGTTAATTTTTCACTCCAGTGC GTCCTGTTGCAGGGACCGGAGCTGATGGGAGCTGCTTCTCCCCATGCCTCACTGGTCCCA GATCANGGCTTCAGGGACAGATGATGAGTCTCAAACGAGCCANCAGGGGTTCTTTTGGGT ATAAATGGGCAATTCGNCCTGTCTTAAGNCTGATGACCTCANCCGTGGTTTTTGGAT Sequence 10054: Found in patent publication WO00/18922 TAGGGCGAATTGGAGCTCCCCGCGTGGCGGCCGAGGTACCTGAGCCAGGAGGAGGCCC GGCCGTGGACCAGGAGCTATTTAACGAATACCAGTTCAGCGTGGACCAACTTATGGAACT GGCCGGCTGAGCTGTGCTACAGCCATCGCCAAGGCATATCCCCCCACGTCCATGTCCAG GAGCCCCCTACTGTCCTGGTCATCTGTGGCCCGGGGAATAATGGAGGAGATGGTCTGGT CTGTGCTCGACACCTCAAACTCTTTGGCTACGAGCCAACCATCTATTACCCCAAAAGGCC

TAACAAGCCCTCTTCACTGCATTGGTGACCCAGTGTCAGAAAATGGACATCCCTTTCCT TGGGGAAATGCCCGCAGAGCCCATGACGATTGATGAACTGTTGAGCTGGTGGTGGATGCC

ATCTTTGGCTTCAGCTTCAAGGGCGATGTTCGGGAAC

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## Table 3-4

Sequence 10047: Found in patent publication WO99/09061 CCCGCGGTGGCCGAGGTACTGCGTGGCCGTCCGCTGTTGTGTGGCCTGGGCCGCAA Α CCGGGCCACGTGGTTGACAATGGGGGAGAAATTGGTAGGACCGTAGAAGCGGATGTGGGG CAGGCAAGCTGAGTACGCCTGGGCAATACCATCCACACCTGAGCAGAAGGGGTTGGTGGG GTTGAAGTTGATGGCAAACTCATGGGAGACCTTCCAGTCTGGGGGTAACTGGGCCCCGAA TCCCAGAGCTGGAAACATCTTATCACTGTCGTAGTCCTGAATGATCTGCCCAACAGCCCA GATGGCCGACAGATATTCGTTGGTGCCCATAGGGTTGATATANTGCAAAGAGGAAGGGTC GAGGGGATTCCCGTTGGAGGCTGTAAAGTCTATTCCAACGGTGAACATGAGCTGGCAGCC TCCCAGGACGTAGTCAAGGAAGGAGTAGT Sequence 10048: Found in patent publication WO00/06698 CCGCGGTGGCGGCCGAGGTCTGAACTGCGGGGTCTCTATCGCACTGCTAGGGGTTCTGCT GCTGGGTGCGCGCGCGCGCGCGGGGCAGAAGCTTTTGAGATTGCTCTGCCACGAGA AAGCAACATTACAGTTCTCATAAAGCTGGGGACCCCGACTCTGCTGGCAAAACCCTGTTA CTTTACCTTTAGCTGCCAGAGTCCTGAGAATCACTTTGTCATANAGATCCAGAAAAATAT TGACTGTATGTCAGGCCCATGTCCTTTTGGGGAGGTTCAGCTTCAGCCCTCGACATCGTT GTTGCCTACCCTCAACAGAACTTTCATCTGGGATGTCAAA Sequence 10049: Found in patent publication WC00/06698 TGATCTTCAGCTGCTCCTGAACTTTGGCACTGTAAGGGTTCTCTGTATCGTATGGTGGTG ACAAAGCTTTTAAAATAACCTNTCCTCCAGTGCATGGAAAAGGACTGGAGTATTTATNAA Sequence 10050: Found in patent publication WC00/06730 ACTIGTGAGCCTGGACTGGCTTCCCCGGCCTCCTACATGAACCCCTTCCCGGTGCTCCAT CTCATCGAGGACTTGAGGCTGGCCTTGGAGATGCTGGAGCTTCCTCAGGAGAGAGCAGCC CTCCTGAGCCAGATCCCTGGCCCAACAGCTGCCTACATAAAGGAATGGTTTGAAGAGAGC TTGTCCCAGGTAAGCCACCACAGTGCTGCTAGTAATGAAACCTACCAGGAACGCTTGGCA CGTCTAGAAGGGGATAAGGAGTCCCTCATATTGCAGGTGAGTGTCCTCACAGACCAAGTA GAAGCCCAGGGAAAAAGATTCGAGACCTGGAAGTGTCTGGAAGGACACCAGGTGAAA CTCAATGCTGCTGAAGAGATGCTTCAACAGGAGCTGCTAAGCCGCACATCTCTTGAGACC CAGAAGCTCGATCTGATGACTGAAGTGTCTGAGCTGAAGCTCAAGCTGGTTGGCATGGAG AAGGAGCAGAGAGCAGGAGAGAAGCAGAGAGAGAGAGTTACTGCAAGAGCTC AGGCCCCTCAAAATCAAAGTGGAAGAGTTTGGAAA Sequence 10051: Found in patent publication WO00/11171 TAATTCTTTGATGGGGTAAATTTGAGCAATTTTCTCGACTTGTCGACATTCGTTATTAAC TGANCAGGAATCAGGANAGGAACCCGGTCCTCTCCACACAGCCCAGCANAGAGCCTACGA CTAGATTTGCATCTTTACGTCCTGCGCGGAGGCTGCTACACACATGCANAAGTCATGCTG GTGGCCTGGACATTGAAGGGAGAGAAGTGGATTTGGGAGACATTTAGGAGGCACCGAAAG CGAAGGAAGCTCCTCCTCCTAAAGCCGAAGCCAAAGCAAAGGCTTTAAAGGCCAAGA Sequence 10052: Found in patent publication WO00/15799 ACCGCGGGGGCCGNGGNTTCNGCCTGGAGGAGCTCAGGGTGGCCGGCATTCACAAG AGGTGGCCCGGACCATCGGCATTTCTGTGGATCCGAGGAGGCGGAACAAGTCCACGGAGT CCCTGCAGACCAACGTGCAGCGGCTGAAGGAGTACCACAAAGGGACCCAAATTCAGCGGT CTGTGCCTACAAACTTCATTAATAACTGCTTGCAGATTGGCAGCTATCTGGTCACTTGAC ATATCCAATGTTGCTATTTTGGTCTGGAGAAAGTTCTCCCTTTCTTCATCTACCTTAATT TCATGTCCATTTTTAAATAATTCAAACATTTTGGGGATGTCACGGCCAATGGAATTTCGA GAAAGCTTTGGATATTTTTTATTCAGTTTCTTCCTTAATCGATTAAGTTCAGGCATTATT TCTGGAACAGCTACGTAAAAGTCTGCAACAATTTCATCATCCCAAATCTTCTGTATCAGA CTAAGTGCCTCCTGCAAATGCAGCTCCATTTTCTTCCGCTATTTTGACCTCTGATGCATT CTCTGTAAATACAGCAACTTTATTGATTTCCGAAGCCAATGGGTATGGCAAACTAANAAC

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ACTGGTAAATGGCTTCACGTTTTTCTTNTTTTCCAGTGCCATATNCAGTGTCAAAATCAA

## **Table 3-4**

A
GGCCGTGGACCAGGAGCTATTTAACGAATACCAGTTCAGCGTGGACCAACTTATGGAACT
GGCCGGGCTGAGCTGTGCTACAGCCATCGCCAAGGCATATCCCCCCACGTCCATGTCCAG

GGCCGGGCTGAGCTGCTACAGCCATCGCCAAGGCATATCCCCCCACGTCCATGTCCAG
GAGCCCCCCTACTGTCCTGGTCATCTGTGGCCCGGGGAATAATGGAGGAGATGGTCTGGT
CTGTGCTCGACACCTCAAACTCTTTGGCTACGAGCCAACCATCTATTACCCCAAAAGGCC
TAACAAGCCCCTCTTCACTGCATTGGTGACCCAGTGTCAGAAAATGGACATCCCTTTCCT
TGGGGAAATGCCCGCAGAGCCCATGACGATTGATGAACTGTTGAGCTGGTGGATGCC
ATCTTTGGCTTCAGCTTCAAGGGCGATGTTCGGGAAC

## Table 3-4

Sequence 10047: Found in patent publication WO99/09061 CCCGCGGTGGCCGAGGTACTGCGTGGCCGTCCGCTGTTGTGTGGCCTGGGCCGCAA

Sequence 10048: Found in patent publication WO00/06698

Sequence 10049: Found in patent publication WO00/0C698

TGATCTTCAGCTGCTCCTGAACTTTGGCACTGTAAGGGTTCTCTGTATCGTATGGTGGTG ACAAAGCTTTTAAAATAACCTNTCCTCCAGTGCATGGAAAAGGACTGGAGTATTTATNAA GT

Sequence 10050: Found in patent publication WO00/06730

Sequence 10051: Found in patent publication WO00/11171

Sequence 10052: Found in patent publication WO00/15799

ACCGCGGGGGCGGCCGNGGNTTCNGCCTGGAGGAGCTCAGGGTGGCCGGCATTCACAAG

A
AGGTGGCCGGACCATCGGCATTTCTGTGGATCCGAGGAGGCGGAACAAGTCCACGGAGT
CCCTGCAGACCAACGTGCAGCGGCTGAAGGAGTACCACAAAGGGACCCAAATTCAGCGGT
CTGTGCCTACAAACTTCATTAATAACTGCTTGCAGATTGGCAGCTATCTGGTCACTTGAC
ATATCCAATGTTGCTATTTTGGTCTGGAGAAAGTTCTCCCTTTCTTCATCTACCTTAATT
TCATGTCCATTTTTAAATAATTCAAACATTTTGGGGATGTCACGGCCAATGGAATTTCGA
GAAAGCTTTGGATATTTTTTATTCAGTTTCTTCCTTAATCGATTAAGTTCAGGCATTATT
TCTGGAACAGC CGTAAAAGTCTGCAACAATTTCATCATCCCAAATCTTCTGTATCAGA
CTAAGTGCCTC GCAAATGCAGCTCCATTTTCTTCCGCTATTTTGACCTCTGATGCATT
CTCTGTAAATACAGCAACTTTATTGATTTCCGAAGCCAATGGGTATGGCAAACTAANAAC
ACTGGTAAATGGCTTCACGTTTTTCTTNTTTTCCAGTGCCATATNCAGTGTCAAAATCAA

## Table 3-4

ANAAAACACTTTGCTTTGGACTAGTTAAAGCCAANAATTTGA Sequence 10053: Found in patent publication WO00/06698 ACATCTTCCAGGACAAGGTAAACTCTGACATGCACTAGGTATGTGCAGATCCCGGCCCT GCCACCCAGCCTCATGCAAGTCATCCCCGACATGACCTTCACGACCGCAATGCAAGGAGG GGAAGAAAGTCACAGCACTGATGAGGACAGCTGCAGAGGTGGCAGTGTGTGGACACAGGA AGTTTGGGCCCCCTCCCTGCCCCAGCTTTCCTAGGCCAGAATTGTGTTTGGCAGTAATTG TCTGTTTAAAAAAATAAAAAGGAGAGGGAAGCGTTCACCGCCACAAATCATAAAATGGACA TGACTGTGGAGTCTTACAGTTCAGGGTTCTTTCATTCACGTCCCTTCCTGTCTCGGTCTG CGGTCTTTACCACATCAATAGGACTTTTTATGCGTCCGGGTTAATTTTTCACTCCAGTGC GTCCTGTTGCAGGGACCGGAGCTGATGGGAGCTGCTTCTCCCCATGCCTCACTGGTCCCA GATCANGGCTTCAGGGACAGATGATGAGTCTCAAACGAGCCANCAGGGGTTCTTTTGGGT ATAAATGGGCAATTCGNCCTGTCTTAAGNCTGATGACCTCANCCGTGGTTTTTGGAT Sequence 10054: Found in patent publication WO00/18922 TAGGGCGAATTGGAGCTCCCCGCGGTGGCGGCCGAGGTACCTGAGCCAGGAGGAGGCCC GGCCGTGGACCAGGAGCTATTTAACGAATACCAGTTCAGCGTGGACCAACTTATGGAACT GGCCGGGCTGAGCTGTGCTACAGCCATCGCCAAGGCATATCCCCCCACGTCCATGTCCAG

GGCCGTGGACCAGGAGCTATTTAACGAATACCAGTTCAGCGTGGACCAACTTATGGAACT GGCCGGGCTGAGCTGTCCTACAGCCATCGCCAAGGCATATCCCCCCACGTCCATGTCCAG GAGCCCCCTACTGTCCTGGTCATCTGTGGCCCGGGGAATAATGGAGGAGATGGTCTGGT CTGTGCTCGACACCTCAAACTCTTTGGCTACGAGCCAACCATCTATTACCCCAAAAGGCC TAACAAGCCCTTCTTCACTGCATTGGTGACCCAGTGTCAGAAAATGGACATCCCTTTCCT TGGGGAAATGCCCGCAGAGCCCATGACGATTGATGAACTGTTGAGCTGGTGGATGCC ATCTTTGGCTTCAGCGTTCAAGGGCGATGTTCGGGAAC

# Table 3-4

Sequence 10047: Found in patent publication WO99/09061 CCCGCGGTGGCCGAGGTACTGCGTGGCCGTCCGCTGTTGTGTGGCCTGGGCCGCAA CCGGGCCACGTGGTTGACAATGGGGGAGAAATTGGTAGGACCGTAGAAGCGGATGTGGGG CAGGCAAGCTGAGTACGCCTGGGCAATACCATCCACCCTGAGCAGAAGGGGTTGGTGGG GTTGAAGTTGATGGCAAACTCATGGGAGACCTTCCAGTCTGGGGGTAACTGGGCCCCGAA TCCCAGAGCTGGAAACATCTTATCACTGTCGTAGTCCTGAATGATCTGCCCAACAGCCCA GATGGCCGACAGATATTCGTTGGTGCCCATAGGGTTGATATANTGCAAAGAGGAAGGGTC GAGGGGATTCCCGTTGGAGGCTGTAAAGTCTATTCCAACGGTGAACATGAGCTGGCAGCC TCCCAGGACGTAGTCAAGGAAGGAGTAGT Sequence 10048: Found in patent publication WO00/06698 CCGCGGTGGCGGCCGAGGTCTGAACTGCGGGGTCTCTATCGCACTGCTAGGGGTTCTGCT GCTGGGTGCGCGCGCGCGCGGGGCAGAAGCTTTTGAGATTGCTCTGCCACGAGA AAGCAACATTACAGTTCTCATAAAGCTGGGGACCCCGACTCTGCTGGCAAAACCCTGTTA CTTTACCTTTAGCTGCCAGAGTCCTGAGAATCACTTTGTCATANAGATCCAGAAAAATAT TGACTGTATGTCAGGCCCATGTCCTTTTGGGGAGGTTCAGCTTCAGCCCTCGACATCGTT GTTGCCTACCCTCAACAGAACTTTCATCTGGGATGTCAAA Sequence 10049: Found in patent publication WO00/06698 TGATCTTCAGCTGCTCCTGAACTTTGGCACTGTAAGGGTTCTCTGTATCGTATGGTGGTG ACAAAGCTTTTAAAATAACCTNTCCTCCAGTGCATGGAAAAGGACTGGAGTATTTATNAA Sequence 10050: Found in patent publication WO00/06730 ACTTGTGAGCCTGGACTGGCTTCCCCGGCCTCCTACATGAACCCCTTCCCGGTGCTCCAT CTCATCGAGGACTTGAGGCTGGCCTTGGAGATGCTGGAGCTTCCTCAGGAGAGAGCAGCC CTCCTGAGCCAGATCCCTGGCCCAACAGCTGCCTACATAAAGGAATGGTTTGAAGAGAGC TTGTCCCAGGTAAGCCACCACAGTGCTGCTAGTAATGAAACCTACCAGGAACGCTTGGCA CGTCTAGAAGGGGATAAGGAGTCCCTCATATTGCAGGTGAGTGTCCTCACAGACCAAGTA GAAGCCCAGGGAGAAAAGATTCGAGACCTGGAAGTGTGTCTGGAAGGACACCAGGTGAAA CTCAATGCTGCAAGAGATGCTTCAACAGGAGCTGCTAAGCCGCACATCTCTTGAGACC CAGAAGCTCGATCTGATGACTGAAGTGTCTGAGCTGAAGCTCAAGCTGGTTGGCATGGAG AAGGAGCAGAGAGCAGGAGGAGAAGCAGAGAGAGAGGAGTTACTGCAAGAGCTC AGGCCCCTCAAAATCAAAGTGGAAGAGTTTGGAAA Sequence 10051: Found in patent publication WO00/11171 GGGAGTGGAGCGTGAACACGTCAGGGGTGGGGGGCGCAGGTCAAGCTTTCACCAGTTTT

Sequence 10052: Found in patent publication WO00/15799
ACCGCGGGGCCGNGGNTTCNGCCTGGAGGAGCTCAGGGTGGCCGGCATTCACAAG

# **Table 3-4**

GGCCGTGGACCAGGAGCTATTTAACGAATACCAGTTCAGCGTGGACCAACTTATGGAACT
GGCCGGGCTGAGCTGCTACAGCCATCGCCAAGGCATATCCCCCCACGTCCATGTCCAG
GAGCCCCCTACTGTCCTGGTCATCTGTGGCCCGGGGAATAATGGAGGAGATGGTCTGGT
CTGTGCTCGACACCTCAAACTCTTTGGCTACGAGCCAACCATCTATTACCCCAAAAGGCC
TAACAAGCCCCTCTTCACTCCATTGGTGACCCAGTGTCAGAAAATGGACATCCCTTTCCT
TGGGGAAATGCCCGCAGAGCCCATGACGATTGATGAACTGTTGAGCTGGTGGTGGATGCC
ATCTTTGGCTTCAGCTTCAAGGGCGATGTTCGGGAAC

#### **Table 3-4**

Sequence 10047: Found in patent publication WO99/09061 CCCGCGGTGGCCGAGGTACTGCGTGGCCGTCCGCTGTTGTGTGGCCTGGGCCGCAA Α CCGGGCCACGTGGTTGACAATGGGGGAGAAATTGGTAGGACCGTAGAAGCGGATGTGGGG CAGGCAAGCTGAGTACGCCTGGGCAATACCATCCACACCTGAGCAGAAGGGGTTGGTGGG GTTGAAGTTGATGGCAAACTCATGGGAGACCTTCCAGTCTGGGGGTAACTGGGCCCCGAA TCCCAGAGCTGGAAACATCTTATCACTGTCGTAGTCCTGAATGATCTGCCCAACAGCCCA GATGGCCGACAGATATTCGTTGGTGCCCATAGGGTTGATATANTGCAAAGAGGAAGGGTC GAGGGGATTCCCGTTGGAGGCTGTAAAGTCTATTCCAACGGTGAACATGAGCTGGCAGCC TCCCAGGACGTAGTCAAGGAAGGAGTAGT Sequence 10048: Found in patent publication WO00/06698 CCGCGGTGGCGGCCGAGGTCTGAACTGCGGGGTCTCTATCGCACTGCTAGGGGTTCTGCT GCTGGGTGCGCGCGCCCCGCGCGGGGCAGAGCTTTTGAGATTGCTCTGCCACGAGA AAGCAACATTACAGTTCTCATAAAGCTGGGGACCCCGACTCTGCTGGCAAAACCCTGTTA CTTTACCTTTAGCTGCCAGAGTCCTGAGAATCACTTTGTCATANAGATCCAGAAAAATAT TGACTGTATGTCAGGCCCATGTCCTTTTGGGGAGGTTCAGCTTCAGCCCTCGACATCGTT GTTGCCTACCCTCAACAGAACTTTCATCTGGGATGTCAAA Sequence 10049: Found in patent publication WQ00/06698 TGATCTTCAGCTGCTCCTGAACTTTGGCACTGTAAGGGTTCTCTGTATCGTATGGTGGTG ACAAAGCTTTTAAAATAACCTNTCCTCCAGTGCATGGAAAAGGACTGGAGTATTTATNAA **GT** Sequence 10050: Found in patent publication WO00/06730 ACTTGTGAGCCTGGACTGCCTCCCGGCCTCCTACATGAACCCCTTCCCGGTGCTCCAT CTCATCGAGGACTTGAGGCTGGCCTTGGAGATGCTGGAGCTTCCTCAGGAGAGAGCAGCC CTCCTGAGCCAGATCCCTGGCCCAACAGCTGCCTACATAAAGGAATGGTTTGAAGAGAGC TTGTCCCAGGTAAGCCACCACAGTGCTGCTAGTAATGAAACCTACCAGGAACGCTTGGCA CGTCTAGAAGGGGTAAGGAGTCCCTCATATTGCAGGTGAGTGTCCTCACAGACCAAGTA GAAGCCCAGGGAAAAGATTCGAGACCTGGAAGTGTCTCTGGAAGGACACCAGGTGAAA CTCAATGCTGCTGAAGAGATGCTTCAACAGGAGCTGCTAAGCCGCACATCTCTTGAGACC CAGAAGCTCGATCTGATGACTGAAGTGTCTGAGCTGAAGCTCAAGCTGGTTGGCATGGAG AAGGAGCAGAGAGCAGGAGAGAAAAGCAGAGGAGTTACTGCAAGAGCTC AGGCCCCTCAAAATCAAAGTGGAAGAGTTTGGAAA Sequence 10051: Found in patent publication WO00/11171 GGGAGTGGAGCGTGAACACGTCAGGGGTGGGGGGCGCAGGTCAAGCTTTCACCAGTTTT TAATTCTTTGATGGGGTAAATTTGAGCAATTTTCTCGACTTGTCGACATTCGTTATTAAC TGANCAGGAATCAGGANAGGAACCCGGTCCTCTCCACACAGCCCAGCANAGAGCCTACGA CTAGATTTGCATCTTTACGTCCTGCGCGGAGGCTGCTACACACATGCANAAGTCATGCTG GTGGCCTGGACATTGAAGGGAGAGAGAGTGGATTTGGGAGACATTTAGGAGGCACCGAAAG CGAAGGAAGCTCCTCCTCCTAAAGCCGAAGCCAAAGCAAAGGCTTTAAAGGCCAAGA Sequence 10052: Found in patent publication WO00/15799 ACCGCGGGGCCGCCGNGGNTTCNGCCTGGAGGAGCTCAGGGTGGCCGGCATTCACAAG AGGTGGCCGGACCATCGGCATTTCTGTGGATCCGAGGAGGCGGAACAAGTCCACGGAGT CCCTGCAGACCAACGTGCAGCGGCTGAAGGAGTACCACAAAGGGACCCAAATTCAGCGGT CTGTGCCTACAAACTTCATTAATAACTGCTTGCAGATTGGCAGCTATCTGGTCACTTGAC

CTGTGCCTACAACTTCATTAATAACTGCTTGCAGATTGGCAGCTATCTGGTCACTTGAC
ATATCCAATGTTGCTATTTTTGGTCTGGAGAAAGTTCTCCCTTTCATCTACCTTAATT
TCATGTCCATTTTTAAATAATTCAAACATTTTGGGATGTCACGGCCAATGGAATTTCGA
GAAAGCTTTGGATATTTTTATTCAGTTTCTTCCTTAATCGACTTAAGTTCAGGCATTATT
TCTGGAACAGCTACGTAAAAGTCTGCAACAATTTCATCCCAAATCTTCTGTATCAGA
CTAAGTGCCTCCTGCAAATGCAGCTCCATTTTCTTCCGCTATTTTTGACCTCTGATGCATT
CTCTGTAAATACAGCAACTTTATTGATTTCCGAAGCCAATGGGTATGGCAAACTAANAAC
ACTGGTAAATGGCTTCACGTTTTTCTTNTTTTCCAGTGCCATATNCAGTGTCAAAAATCAA

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Α

## Table 3-4

ANAAACACTTTGCTTTGGACTAGTTAAAGCCAANAATTTGA
Sequence 10053: Found in patent publication W000/06698
ACATCTTCCAGGACAAGGTAAACTCTGACATGCACTAGGTATGTGCAGATCCCGGCCCT
GCCACCCAGCCTCATGCAAGTCATCCCCGACATGCACTCACGACCGCAATGCAAGGAGG
GGAAGAAAGTCACAGCACTGATGAGGACAGCTGCAGAGGTGGCAGTGTGTGGACACAGGA
AGTTTGGGCCCCCTCCCTGCCCCAGCTTTCCTAGGCCAGAATTGTTTTGGCAGTAATTG
TCTGTTTAAAAAAATAAAAAGGAGAGAGGAAGCGTTCACCGCCACAAATCATAAAATGGACA
TGACTGTGGAGTCTTACAGTTCAGGGTTCTTTCATTCACGTCCCTTCCTGTCTCGGTCTG
CGGTCTTTACCACATCAATAGGACTTTTTATGCGTCCGGGTTAATTTTTCACTCCAGTGC
GTCCTGTTGCAGGGACCGGAGCTGATGGGAGCTGCTTCTCCCCATGCCTCACTGGTCCCA
GATCANGGCTTCAGGGACAGATGATGAGTCTCAAACGAGCCANCAGGGGTTCTTTTGGGT
ATAAATGGGCAATTCGNCCTGTCTTAAGNCTGATGACCTCANCCGTGGTTTTTTGGAT
Sequence 10054: Found in patent publication W000/18922
TAGGGCGAATTGGAGCTCCCCGCGGTGGCGGCCGAGGTACCTGAGCCAGGAGGAGGCCC

GGCCGTGGACCAGGAGCTATTTAACGAATACCAGTTCAGCGTGGACCAACTTATGGAACT GGCCGGGCTGAGCTGCTACAGCCATCGCCAAGGCATATCCCCCCACGTCCATGTCCAG GAGCCCCCTACTGTCCTGGTCATCTGTGGCCCGGGGAATAATGGAGGAGATGGTCTGGT CTGTGCTCGACACCTCAAACTCTTTGGCTACGAGCCAACCATCTATTACCCCAAAAGGCC TAACAAGCCCCTCTTCACTGCATTGGTGACCAGTGTCAGAAAATGGACATCCCTTTCCT TGGGGAAATGCCCGCAGAGCCCATGACGATTGATGAACTGTTGAGCTGGTGGATGCC ATCTTTGGCTTCAGCTTCAAGGGCGATGTTCGGGAAC

# Table 3-4

Sequence 10047: Found in patent publication WO99/09061
CCCGCGGTGGCCGAGGTACTGCGTGGCCGTCCGCTGTTGTGTGGCCTGGGCCGCAA
A

CCGGGCCACGTGGTTGACAATGGGGGAGAAATTGGTAGGACCGTAGAAGCGGATGTGGGG CAGGCAAGCTGAGTACGCCTGGGCAATACCATCCACACCTGAGCAGAAGGGGTTGGTGGG GTTGAAGTTGATGGCAAACTCATGGGAGACCTTCCAGTCTGGGGGTAACTGGGCCCCGAA TCCCAGAGCTGGAAACATCTTATCACTGTCGTAGTCCTGAATGATCTGCCCAACAGCCCA GATGGCCGACAGATATTCGTTGGTGCCCATAGGGTTGATATANTGCAAAGAGGAAGGGTC GAGGGGATTCCCGTTGGAGGCTGTAAAGTCTATTCCAACGGTGAACATGAGCTGGCAGCC TCCCAGGACGTAGTCAAGGAAGGAGTAGT

Sequence 10048: Found in patent publication WO00/06698

CCGCGGTGGCGGCCGAGGTCTGAACTGCGGGGTCTCTATCGCACTGCTAGGGGTTCTGCT GCTGGGTGCGGCGCCCGCGCGCGGGGGCAGAAGCTTTTGAGATTGCTCTGCCACGAGA AAGCAACATTACAGTTCTCATAAAGCTGGGGACCCCGACTCTGCTGGCAAAACCCTGTTA CATCGTCTTTTCTAAAAGACATATAACCATGTTGTCCATCAAGTCTGGAGAAAAGAATAGT CTTTACCTTTAGCTGCCAGAGTCCTGAGAATCACTTTGTCATANAGATCCAGAAAAATAT TGACTGTATGTCAGGCCCATGTCCTTTTGGGGAGGTTCAGCTTCAGCCCTCGACATCGTT GTTGCCTACCCTCAACAGAACTTTCATCTGGGATGTCAAA

Sequence 10049: Found in patent publication WO00/06698

TGATCTTCAGCTGCTCCTGAACTTTGGCACTGTAAGGGTTCTCTGTATCGTATGGTGGTG ACAAAGCTTTTAAAATAACCTNTCCTCCAGTGCATGGAAAAGGACTGGAGTATTTATNAA GT

Sequence 10050: Found in patent publication WO00/06730

ACTTGTGAGCCTGGACTGCTTCCCCGGCCTCCTACATGAACCCCTTCCCGGTGCTCCAT CTCATCGAGGACTTGAGGCTGGCCTTGGAGATGCTGGAGCTTCCTCAGGAGAGAGCAGCC CTCCTGAGCCAGATCCCTGGCCCAACAGCTGCCTACATAAAGGAATGGTTTGAAGAGAGC TTGTCCCAGGTAAGCCACCACAGTGCTGCTAGTAATGAAACCTACCAGGAACGCTTGGCA CGTCTAGAAGGGGATAAGGAGTCCCTCATATTGCAGGTGAGTGTCCTCACAGACCAAGTA GAAGCCCAGGGAGAAAAGATTCGAGACCTGGAAGTGTGTCTGGAAGGACACCAGGTGAAA CTCAATGCTGCTGAAGAGATGCTTCAACAGGAGCTGCTAAGCCGCACATCTCTTGAGACC CAGAAGCTCGATCTGATGACTGAAGTGTCTGAGCTCAAGCTGGTTGGCATGGAG AAGGAGCAGAGAGAGCAGGAGAAAAGCAGAGGAGTTACTGCAAGAGCTC AGGCCCCTCAAAATCAAAGTGGAAGAGTTTGGAAA

Sequence 10051: Found in patent publication WO00/11171

Sequence 10052: Found in patent publication WO00/15799

ACCGCGGGGGCCGNGGNTTCNGCCTGGAGGAGCTCAGGGTGGCCGGCATTCACAAG

A AGGTGGCCCGGACCATCGGCATTTCTGTGGATCCGAGGAGGCGGAACAAGTCCACGGAGT CCCTGCAGACCAACGTGCAGCGGCTGAAGGAGTACCACAAAGGGACCCAAATTCAGCGGT CCTGCAGACCAAACTTCATTAATAACTGCTTGCAGATTGGCAGCTATCTGGTCACTTGAC ATATCCAATGTTGCTATTTTGGTCTGGAGAAAGTTCTCCCTTTCTTCATCTACCTTAATT TCATGTCCATTTTTAAATAATTCAAACATTTTTGGGGATGTCACGGCCAATGGAATTTCGA GAAAGCTTTGGATATTTTTATTCAGTTTCTTCCTTAATCGATTAAGTTCAGGCATTATT TCTGGAACAGCTACGTAAAAGTCTGCAACAATTTCATCATCCCAAATCTTCTGTATCAGA CTAAGTGCCTCCTGCAAATGCAGCATTTCTTCCGCTATTTTGACCTCTGATGCATT CTCTGTAAATACAGCAACTTTATTGATTTCCGAAGCCAATGGGTATGGCAAACTAANAAC ACTGGTAAATGCGCTTCACGTTTTTCTTNTTTTCCAGTGCCATATNCAGTGTCAAAATCAA

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## Table 3-4

ANAAAACACTTTGCTTTGGACTAGTTAAAGCCAANAATTTGA
Secuence 10053: Found in patent publication WO00/06698
ACATCTTCCAGGACAAGGTAAACTCTGACATGCACTAGGTATGTGCAGATCCCGGCCCCT
GCCACCCAGCCTCATGCAAGTCATCCCCGACATGACCTTCACGACCGCAATGCAAGGAGG
GGAAGAAAGTCACAGCACTGATGAGGACAGCTGCAGAGGTGGCAGTGTGGACACAGGA
AGTTTGGGCCCCCTCCCTGCCCCAGCTTTCCTAGGCCAGAATTGTTTTGGCAGTAATTG
TCTGTTTAAAAAAAATAAAAAGGAGAGAGGAAGCGTTCACCGCCACAAATCATAAAATGGACA
TGACTGTGGAGTCTTACAGTTCAGGGTTCTTTCATTCACGTCCCTTCCTGTCTCGGTCTG
CGGTCTTTACCACATCAATAGGACTTTTTTATGCGTCCGGGTTAATTTTTCACTCCAGTGC
GTCCTGTTGCAGGGACCGGAGCTGATGGGAGCTGCTTCTCCCCATGCCTCACTGGTCCCA
GATCANGGCTTCAGGGACAGATGATGAGTCTCAAACGAGCCANCAGGGGTTCTTTTGGGT
ATAAATGGGCAATTCGNCCTGTCTTAAGNCTGATGACCTCANCCGTGGTTTTTTGGAT
Sequence 10054: Found in patent publication WO00/18922
TAGGGCGAATTGGAGCTCCCCGCGGTGGCGGCCGAGGTACCTGAGCCAGGAGGAGGCCC

GGCCGTGGACCAGGAGCTATTTAACGAATACCAGTTCAGCGTGGACCAACTTATGGAACT GGCCGGGCTGAGCTGTCACAGCCATCGCCAAGGCATATCCCCCCACGTCCATGTCCAG GAGCCCCCTACTGTCCTGGTCATCTGTGGCCCGGGGAATAATGGAGGAGATGGTCTGGT CTGTGCTCGACACCTCAAACTCTTTGGCTACGAGCCAACCATCTATTACCCCAAAAGGCC TAACAAGCCCCTCTTCACTGCATTGGTGACCCAGTGTCAGAAAATGGACATCCCTTTCCT TGGGGAAATGCCCGCAGAGCCCATGACGATTGATGAACTGTTGAGCTGGTGGTGGATGCC ATCTTTGGCTTCAACGTCCAAGGGCGATGTTCGGGAAC

# Table 3-4

Sequence 10047: Found in patent publication WO99/09061
CCCGCGGTGGCCGAGGTACTGCGTGGCCGTCGCTGTTGTGTGGCCTGGGCCGCAA
A
CCGGGCCACGTGGTTGACAATGGGGGAGAAATTGGTAGGACCGTAGAAGCGGATGTGGGG

Sequence 10048: Found in patent publication WO00/06698

CCGCGTGGCGGCCGAGGTCTGAACTGCGGGGTCTCTATCGCACTGCTAGGGGTTCTGCT GCTGGGTGCGGCGCCCTGCCGCGGGGGCAGAAGCTTTTGAGATTGCTCTGCCACGAGA AAGCAACATTACAGTTCTCATAAAGCTGGGGACCCCGACTCTGCTGGCAAAACCCTGTTA CATCGTCTTTTCTAAAAGACATATAACCATGTTGTCCATCAAGTCTGGAGAAAAGAATAGT CTTTACCTTTAGCTGCCAGAGTCCTGAGAATCACTTTGTCATANAGATCCAGAAAAATAT TGACTGTATGTCAGGCCCATGTCCTTTTGGGGAGGTTCAGCTTCAGCCCTCGACATCGTT GTTGCCTACCCTCAACAGAACTTTCATCTGGGATGTCAAA

Sequence 10049: Found in patent publication WO00/06698

TGATCTTCAGCTGCTCCTGAACTTTGGCACTGTAAGGGTTCTCTGTATCGTATGGTGGTG ACAAAGCTTTTAAAATAACCTNTCCTCCAGTGCATGGAAAAGGACTGGAGTATTTATNAA GT

Sequence 10050: Found in patent publication WO00/06730

ACTTGTGAGCCTGGACTGGCTTCCCCGGCCTCCTACATGAACCCCTTCCCGGTGCTCCAT
CTCATCGAGGACTTGAGGCTGGCCTTGGAGATGCTGGAGCTTCCTCAGGAGAGAGCAGCC
CTCCTGAGCCAGATCCCTGGCCCAACAGCTGCCTACATAAAGGAATGGTTTGAAGAGAGC
TTGTCCCAGGTAAGCCACCACAGTGCTGCTAGTAATGAAACCTACCAGGAACGCTTGGCA
CGTCTAGAAGGGGATAAGGAGTCCCTCATATTGCAGGTGAGTGTCCTCACAGACCAAGTA
GAAGCCCAGGGAGAAAAGATTCGAGACCTGGAAGTGTCTCTGAAAGGACCCAGGTGAAA
CTCAATGCTGCTGAAGAGTTCTCAACAGGAGCTGCTAAGCCGCACATCTCTTGAGACC
CAGAAGCTCGATCTGATGACTGAAGTGTCTGAGCTCAAGCTCGATTGGCATGAG
AAGGAGCAGAGAGAGAGAGAGAAAAGCAGAGGAGTTACTGCAAGAGCTC
AGGCCCCTCAAAATCAAAGTGGAAGAGTTTGGAAA

Sequence 10051: Found in patent publication WO00/11171

Sequence 10052: Found in patent publication WO00/15799

ACCGCGGGGCGGCCGNGGNTTCNGCCTGGAGGAGCTCAGGGTGGCCGGCATTCACAAG

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## **Table 3-4**

ANAAAACACTTTGCTTTGGACTAGTTAAAGCCAANAATTTGA Sequence 10053: Found in patent publication WC00/06698 ACATCTTCCAGGACAAGGTAAACTCTGACATGCACTAGGTATGTGCAGATCCCGGCCCCT GCCACCCAGCCTCATGCAAGTCATCCCCGACATGACCTTCACGACCGCAATGCAAGGAGG GGAAGAAGTCACAGCACTGATGAGGACAGCTGCAGAGGTGGCAGTGTGTGGACACAGGA AGTTTGGGCCCCCTCCCTGCCCCAGCTTTCCTAGGCCAGAATTGTGTTTGGCAGTAATTG TCTGTTTAAAAAAATAAAAAGGAGGAGGAAGCGTTCACCGCCACAAATCATAAAATGGACA TGACTGTGGAGTCTTACAGTTCAGGGTTCTTTCATTCACGTCCCTTCCTGTCTCGGTCTG CGGTCTTTACCACATCAATAGGACTTTTTATGCGTCCGGGTTAATTTTTCACTCCAGTGC GTCCTGTTGCAGGGACCGGAGCTGATGGGAGCTGCTTCTCCCCATGCCTCACTGGTCCCA GATCANGGCTTCAGGGACAGATGATGAGTCTCAAACGAGCCANCAGGGGTTCTTTTGGGT ATAAATGGGCAATTCGNCCTGTCTTAAGNCTGATGACCTCANCCGTGGTTTTTGGAT Sequence 10054: Found in patent publication WO00/18922 TAGGGCGAATTGGAGCTCCCCGCGTGGCGGCCGAGGTACCTGAGCCAGGAGGAGGCCC GGCCGTGGACCAGGAGCTATTTAACGAATACCAGTTCAGCGTGGACCAACTTATGGAACT GGCCGGCTGAGCTGTGCTACAGCCATCGCCAAGGCATATCCCCCCACGTCCATGTCCAG GAGCCCCCCTACTGTCCTGGTCATCTGTGGCCCGGGGAATAATGGAGGAGATGGTCTGGT CTGTGCTCGACACCTCAAACTCTTTGGCTACGAGCCAACCATCTATTACCCCAAAAGGCC

TAACAAGCCCTCTTCACTGCATTGGTGACCCAGTGTCAGAAAATGGACATCCCTTTCCT TGGGGAAATGCCCGCAGAGCCCATGACGATTGATGAACTGTTGAGCTGGTGGTGGATGCC

ATCTTTGGCTTCAGCTTCAAGGGCGATGTTCGGGAAC

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<u>Table 3-5</u>

C	Ann No	Database hit	GI Nbr
Sequence #	Acc No		345130
Sequence 10055	A02759	Genbank	2293812
Sequence 10056	A36501	Genbank	
Sequence 10057	A38056	Genbank	2294706
Sequence 10058	A46722	Genbank	2300822
Sequence 10059	A98295	Genbank	6781400
Sequence 10060	AB000584	Genbank	1813326
Sequence 10061	AB000888	Genbank	2467297
Sequence 10062	AB <b>002323</b>	Genbank	2224590
Sequence 10063	AB0 <b>02337</b>	Genbank	6683125
Sequence 10064	AB0 <b>02383</b>	Genbank	2224710
Sequence 10065	AB002533	Genbank	1944124
Sequence 10066	AB003102	Genbank	1945608
Sequence 10067	AB00 <b>4064</b>	Genbank	6983841
Sequence 10068	AB0 <b>04885</b>	Genbank	2217932
Sequence 10069	AB006534	Genbank	2924619
Sequence 10070	AB007170	Genbank	3077765
Sequence 10071	AB007956	Genbank	3413930
Sequence 10072	AB007964	Genbank	3413939
Sequence 10073	AB008226	Genbank	3370992
Sequence 10074	AB011004	Genbank	3273315
Sequence 10075	AB011083	Genbank	3043545
Sequence 10076	AB011100	Genbank	6683714
Sequence 10077	AB011120	Genbank	3043619
Sequence 10078	AB011165	Genbank	3043709
Sequence 10079	AB011399	Genbank	3452571
Sequence 10080	AB014511	Genbank	3327035
Sequence 10081	AB014570	Genbank	3327153
Sequence 10082	AB014579	Genbank	3327171
Sequence 10083	AB014583	Genbank	6635130
Sequence 10084	AB015639	Genbank	5821139
Sequence 10085	AB015798	Genbank	6681591
Sequence 10086	AB016089	Genbank	5821146
Sequence 10087	AB018257	Genbank	3882148
Sequence 10088	AB018274	Genbank	3882182
Sequence 10089	AB018284	Genbank	3882202
Sequence 10090	AB018304	Genbank	3882242
Sequence 10091	AB018330	Genbank	3882294
Sequence 10092	AB018340	Genbank	3882314
Sequence 10093	AB019564	Genbank	3885367
Sequence 10094	AB019568	Genbank	3885371
Sequence 10095	AB020335	Genbank	6518494
Seguence 10096	AB020676	Genbank	4240226
Sequence 10097	AB020683	Genbank	4240240
Sequence 10098	AB020694	Genbank	4240262
Sequence 10099	AB020864	Genbank	4003384
Sequence 10100	AB020867	Genbank	4003387
Sequence 10101	AB021179	Genbank	4062855
Sequence 10102	AB021467	Genbank	4586518
Sequence 10103	AB <b>021663</b>	Genbank	4996450
Sequence 10104	AB023050	Genbank	56 <b>72605</b>
Sequence 10105	AB023051	Genbank	5672606
Sequence 10106	AB023140	Genbank	4589477
Sequence 10107	AB023208	Genbank	4589625
Sequence 10108	AB024334	Genbank	601 <b>6837</b>
Sequence 10109	AB025355	Genbank	5103146

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<u>Table 3-5</u>

Sequence 10110	AB025959	Genbank	6520072
Sequence 10111	AB026894	Genbank	5821423
Sequence 10112	AB027466	Genbank	6172220
Sequence 10113	AB028624	Genbank	5103045
Sequence 10114	AB028859	Genbank	6567165
Sequence 10115	AB028893	Genbank	6552364
Sequence 10116	AB028975	Genbank	5689440
Sequence 10117	AB029001	Genbank	5689492
Sequence 10118	AB029025	Genbank	5689540
Sequence 10119	AB031052	Genbank	6978947
Sequence 10120	AB032957	Genbank	
Sequence 10121	AB032959	Genbank	6329818
Sequence 10121	AB032982	Genbank	6329832
Sequence 10123	AB033017	Genbank	6330122
Sequence 10123	AB033050	Genbank	6330344
Sequence 10125	AB033074		6330623
Sequence 10125	AB033074 AB033079	Genbank	6330846
Sequence 10127		Genbank	6382025
•	AB034205	Genbank	6899845
Sequence 10128	AB034206	Genbank	6177737
Sequence 10129	AB035207	Genbank	6469033
Sequence 10130	AB037740	Genbank	7243018
Sequence 10131	AB037763	Genbank	7243064
Sequence 10132	AB037767	Genbank	7243072
Sequence 10133	AB037817	Genbank	7243172
Sequence 10134	AB037850	Genbank	7243238
Sequence 10135	AC000049	Genbank	5705995
Sequence 10136	AC000064	Genbank	1669369
Sequence 10137	AC000080	Genbank	7123907
Sequence 10138	AC000120	Genbank	1809224
Sequence 10139	AC000353	Genbank	6970735
Sequence 10140	AC000378	Genbank	2270906
Sequence 10141	AC000394	Genbank	2133911
Sequence 10142	AC001226	Genbank	2133862
Sequence 10143	AC001642	Genbank	1945597
Sequence 10144	AC002038	Genbank	2226439
Sequence 10145	AC002039	Genbank	2342716
Sequence 10146	AC002051	Genbank	7798752
Sequence 10147	AC002064	Genbank	2076723
Sequence 10148	AC002347	Genbank	2828783
Sequence 10149	AC002350	Genbank	2961443
Sequence 10150	AC002357	Genbank	2815516
Sequence 10151	AC002400	Genbank	2576344
Sequence 10152	AC002470	Genbank	6478944
Sequence 10153	AC002528	Genbank	2388554
Sequence 10154	AC002531	Genbank	2580476
Sequence 10155	AC002544	Genbank	3337382
Sequence 10156	AC002553	Genbank	3126783
Sequence 10157	AC002984	Genbank	2443872
Sequence 10158	AC003004	Genbank	2522284
Sequence 10159	AC003007	Genbank	2911728
Sequence 10160	AC003012	Genbank	2547258
Sequence 10161	AC003025	Genbank	3337308
Sequence 10162	AC003047	Genbank	31 <b>69149</b>
Sequence 10163	AC003103	Genbank	2842782
Sequence 10164	AC003111	Genbank	2636670
Sequence 10165	AC003663	Genbank	3 <b>097871</b>

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<u>Table 3-5</u>

Sequence 10166	AC003688	Genbank	3789719
Sequence 10167	AC003692	Genbank	2731601
Sequence 10168	AC003973	Genbank	2739352
	AC004008		-
Sequence 10169		Genbank	2781388
Sequence 10170	AC004036	Genbank	2811104
Sequence 10171	AC004041	Genbank	2811099
Sequence 10172	AC004067	Genbank	3851208
Sequence 10173	AC004084	Genbank	2822156
Sequence 10174	AC004111	Genbank <sup>-</sup>	2828786
Sequence 10175	AC004148	Genbank	3482960
Sequence 10176	AC004151	Genbank	2896795
Sequence 10177	AC004213	Genbank	3386628
Sequence 10178	AC004218	Genbank	4263838
		Genbank	
Sequence 10179	AC004236		2914668
Sequence 10180	AC004258	Genbank	2924759
Sequence 10181	AC004386	Genbank	3046272
Sequence 10182	AC004388	Genbank	3046271
Sequence 10183	AC004453	Genbank	2979600
Sequence 10184	AC004496	Genbank	2996645
Sequence 10185	AC004527	Genbank	4760422
Sequence 10186	AC004530	Genbank	3108050
Sequence 10187	AC004551	Genbank	3309005
Sequence 10188	AC004582	Genbank	3337307
Sequence 10189	AC004584	Genbank	3417305
Sequence 10190	AC004596	Genbank	4210521
Sequence 10191	AC004637	Genbank	3094998
		Genbank	3115345
Sequence 10192	AC004668	Genbank	
Sequence 10193	AC004682		3337388
Sequence 10194	AC004687	Genbank	3258613
Sequence 10195	AC004707	Genbank	3249127
Sequence 10196	AC004742	Genbank	3152630
Sequence 10197	AC004757	Genbank	3 <b>341707</b>
Sequence 10198	AC004810	Genbank	3 <b>264567</b>
Sequence 10199	AC004816	Genbank	4156209
Sequence 10200	AC004821	Genbank	4753291
Sequence 10201	AC004832	Genbank	6624129
Sequence 10202	AC004835	Genbank	450 <b>8154</b>
Sequence 10203	AC004854	Genbank	4827328
Sequence 10204	AC004865	Genbank	4156195
Sequence 10205	AC004878	Genbank	3213119
Sequence 10206	AC004884	Genbank	4156189
Sequence 10207	AC004904	Genbank	4156180
	AC004904 AC004908		
Sequence 10208		Genbank	4156179
Sequence 10209	AC004913	Genbank	3213084
Sequence 10210	AC004918	Genbank	4153871
Sequence 10211	AC004935	Genbank	4156170
Sequence 10212	AC004938	Genbank	3213059
Sequence 10213	AC004943	Genbank	3924 <b>671</b>
Sequence 10214	AC004955	Genbank	4753 <b>267</b>
Sequence 10215	AC004968	Genbank	4156160
Sequence 10216	AC004969	Genbank	41561 <b>59</b>
Sequence 10217	AC004982	Genbank	3419846
Sequence 10218	AC004983	Genbank	4309885
Sequence 10219	AC004985	Genbank	5708490
Sequence 10213	AC004999	Genbank	3970 <b>963</b>
	AC005023	Genbank	3900847
Sequence 10221	AC000023	Genbank	390004/

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Sequence 10222	AC005035	Genbank	3953506
Sequence 10223	AC005041	Genbank	4508118
Sequence 10224	AC005057	Genbank	6587915
Sequence 10225	AC005058	Genbank	4156135
Sequence 10226	AC005069	Genbank	
Sequence 10227			4753224
	AC005070	Genbank	3406049
Sequence 10228	AC005071	Genbank	4508112
Secuence 10229	AC005095	Genbank	4753220
Secuence 10230	AC005101	Genbank	4508108
Sequence 10231	AC005154	Genbank	3242763
Secuence 10232	AC005180	Genbank	36 <b>87281</b>
Secuence 10233	AC005189	Genbank	3264580
Secuence 10234	AC005193	Genbank	3309101
Sequence 10235	AC005203	Genbank	3273381
Sequence 10236	AC005212	Genbank	3287444
Sequence 10237	AC005218	Genbank	3282162
Sequence 10238	AC005258	Genbank	3289992
Sequence 10239	AC005263	Genbank	3289978
Sequence 10240	ACC05300	Genbank	
Sequence 10241	ACC05316		6492490
Sequence 10247		Genbank	3738342
	AC005318	Genbank	3885345
Sequence 10243	AC005325	Genbank	3366581
Sequence 10244	AC005343	Genbank	4165008
Sequence 10245	AC005346	Genbank	3366566
Sequence 10246	AC005368	Genbank	3367506
Sequence 10247	AC005378	Genbank	4508139
Sequence 10248	AC005412	Genbank	5931398
Sequence 10249	AC005480	Genbank	4835818
Sequence 10250	AC005488	Genbank	58361 <b>94</b>
Sequence 10251	AC005520	Genbank	5001541
Sequence 10252	AC005531	Genbank	4153875
Sequence 10253	AC005550	Genbank	34786 <b>68</b>
Sequence 10254	AC005606	Genbank	3540172
Sequence 10255	AC005612	Genbank	3540153
Sequence 10256	AC005618	Genbank	3548785
Sequence 10257	AC005630	Genbank	4159882
Sequence 10258	AC005661	Genbank	3818348
Sequence 10259	AC005740	Genbank	3687210
Sequence 10260	AC005754	Genbank	3688074
Sequence 10261	AC005786	Genbank	3702287
Sequence 10262	AC005789	Genbank	3702281
Sequence 10263	AC005799	Genbank	3789715
Sequence 10264	AC005828	Genbank	3789713
Sequence 10265	AC005876	Genbank	6249672
Sequence 10266	AC005971	Genbank	4314421
Sequence 10267	AC006001	Genbank	5708496
Sequence 10268	AC006008	Genbank	5091655
Sequence 10269	AC006017	Genbank	4508141
Sequence 10270	AC006040	Genbank	4314426
Sequence 10271	AC006055	Genbank	4071011
Sequence 10272	AC006057	Genbank	4731048
Sequence 10273	AC006059	Genbank	4544348
Sequence 10274	AC006064	Genbank	4572650
Sequence 10275	AC006116	Genbank	3962497
Sequence 10276	AC006160	Genbank	
Sequence 10277	AC006160 AC006254		5701616
00436108 10217	AC000254	Genbank	4726099

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Sequence 10278	AC006255	Genbank	5306223
Sequence 10279	AC006257	Genbank	4092478
Sequence 10280	AC006261	Genbank	4079613
Sequence 10281	AC006312	Genbank	4582483
Sequence 10282	AC006347	Genbank	4309919
	_		
Sequence 10283	AC006389	Genbank	4309883
Sequence 10284	AC006445	Genbank	6721147
Sequence 10285	AC006454	Genbank	5732182
Sequence 10286	AC006461	Genbank	4508128
Sequence 10287	AC006477	Genbank	7243871
Sequence 10288	AC006485	Genbank	4206211
Sequence 10289	AC006487	Genbank	4314418
Sequence 10290	AC006501	Genbank	4309874
Sequence 10291	AC006504	Genbank	4220442
Sequence 10292	AC006509	Genbank	5815507
Sequence 10293	AC006518	Genbank	4713939
Sequence 10293	AC006543	Genbank	4388738
		Genbank	4558534
Sequence 10295	AC006561		
Sequence 10296	AC006581	Genbank	4914350
Sequence 10297	AC006960	Genbank	4337211
Sequence 10298	AC007041	Genbank	5708471
Sequence 10299	AC007055	Genbank	4885691
Sequence 10300	AC007191	Genbank	4558643
Sequence 10301	AC007204	Genbank	4559317
Sequence 10302	AC007225	Genbank	6715703
Sequence 10303	AC007226	Genbank	6715702
Sequence 10304	AC007227	Genbank	6456148
Sequence 10305	AC007325	Genbank	7229741
Sequence 10306	AC007528	Genbank	5263309
Sequence 10307	AC007546	Genbank	5668756
Sequence 10308	AC007637	Genbank	8439781
Sequence 10309	AC007684	Genbank	5836173
Sequence 10310	AC007708	Genbank	5923689
Sequence 10311	AC007790	Genbank	5042474
Sequence 10317	AC007750 AC007842	Genbank	5080755
•	AC007967	Genbank	5931455
Sequence 10313		Genbank	6143846
Sequence 10314	AC008082		6970743
Sequence 10315	AC008101	Genbank	
Sequence 10316	AC008116	Genbank	6001959
Sequence 10317	AC008124	Genbank	5815491
Sequence 10318	AC008179	Genbank	5931440
Sequence 10319	AC008269	Genbank	7243890
Sequence 10320	AC00B498	Genbank	6850299
Sequence 10321	AC008893	Genbank	7259691
Sequence 10322	AC009363	Genbank	6648145
Sequence 10323	AC009509	Genbank	6492473
Sequence 10324	AC009946	Genbank	6604542
Sequence 10325	AC011331	Genbank	9929686
Sequence 10326	AC011362	Genbank	6094545
Sequence 10327	AC015853	Genbank	6721261
Sequence 10328	AC020663	Genbank	6682593
Sequence 10329	AD000092	Genbank	1905905
Sequence 10330	AE000658	Genbank	2358019
	AF000160	Genbank	2306783
Sequence 10331	AF000160 AF001437	Genbank	2316039
Sequence 10332			3355302
Sequence 10333	AF001549	Genbank	3333302

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<u>Table 3-5</u>

Sequence 10334	AF0C3001	Genbank	2529443
Sequence 10335	AF005888	Genbank	2738487
Sequence 10336	AF0C6043	Genbank	2674061
Sequence 10337	AF0C6088	Genbank	2282041
Sequence 10338	AF006501	Genbank	4836840
Sequence 10339	AF007140	Genbank	2852616
Sequence 10340	AF0C7544	Genbank	2970122
Sequence 10341	AF010238	Genbank	2282063
Sequence 10342	AF011390	Genbank	3298567
Sequence 10343	AF012072	Genbank	9967556
Sequence 10344	AF014807	Genbank	2338731
Sequence 10345	AF014955	Genbank	2772828
Sequence 10346	AF015812	Genbank	2599359
Sequence 10347	AF020351	Genbank	
Sequence 10348	AF020352	Genbank	2655052
Sequence 10349	AF021819	Genbank	2655054
Sequence 10349	AF022211	Genbank	2460317
Sequence 10350	AF026126		2465723
Sequence 10352	AF026126	Genbank	2828601
Sequence 10352	AF028832	Genbank	2559009
_ '	AF028632 AF029786	Genbank	3287488
Sequence 10354 Sequence 10355	AF029786 AF031416	Genbank	3403166
		Genbank	3213216
Sequence 10356	AF032387	Genbank	2641556
Sequence 10357	AF033095	Genbank	2645728
Sequence 10358	AF034759	Genbank	2653648
Sequence 10359	AF035286	Genbank	2661038
Sequence 10360	AF035295	Genbank	2661050
Sequence 10361	AF035313	Genbank	2661075
Sequence 10362	AF037222	Genbank	2921499
Sequence 10363	AF037447	Genbank	6466790
Sequence 10364	AF037448	Genbank	3037012
Sequence 10365	AF038196	Genbank	2795917
Sequence 10366	AF038200	Genbank	2795921
Sequence 10367	AF038554	Genbank	28955 <b>58</b>
Sequence 10368	AF038954	Genbank	3329377
Sequence 10369	AF038955	Genbank	332937 <b>9</b>
Sequence 10370	AF038958	Genbank	3329385
Sequence 10371	AF038966	Genbank	2791681
Sequence 10372	AF039029	Genbank	3834389
Sequence 10373	AF042836	Genbank	3776241
Sequence 10374	AF042857	Genbank	2952271
Sequence 10375	AF044956	Genbank	5326827
Sequence 10376	AF044957	Genbank	4164445
Sequence 10377	AF044959	Genbank	3348136
Sequence 10378	AF047181	Genbank	2 <b>9</b> 09 <b>853</b>
Sequence 10379	AF047431	Genbank	5514630
Sequence 10380	AF047436	Genbank	3335127
Sequence 10381	AF047439	Genbank	3 <b>33</b> 51 <b>33</b>
Sequence 10382	AF047440	Genbank	3 <b>33</b> 51 <b>35</b>
Sequence 10383	AF048686	Genbank	573769 <b>8</b>
Sequence 10384	AF049140	Genbank	294730 <b>0</b>
Sequence 10385	AFC49613	Genbank	332942 <b>8</b>
Sequence 10386	AF050637	Genbank	4164451
Sequence 10387	AF050638	Genbank	5 <b>3</b> 26 <b>819</b>
Sequence 10388	AF050639	Genbank	41644 <b>53</b>
Sequence 10389	AFC51894	Genbank	309511 <b>0</b>

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Sequence 10390	AF052101	Genbank	33604 <b>08</b>
Sequence 10391		Genbank	336043 <b>8</b>
Sequence 10392		Genbank	2981453
Sequence 10393		Genbank	3135305
Sequence 10394		Genbank	3283048
Sequence 1039		Genbank	6670884
Sequence 10396		Genbank	4092059
Sequence 10397		Genbank	3005703
Sequence 10398		Genbank	3005711
Sequence 10399		Genbank	30057 <b>59</b>
Sequence 1039		Genbank	3335 1 3
		Genbank	<b>3</b> 500. )
Sequence 1040		Genbank	<b>328</b> 30
Sequence 1040	2 AF063020		334785
Sequence 1040		Genbank	3152667
Sequence 1040	4 AF064607	Genbank	
Sequence 1040	5 AF067518	Genbank	3850315
Sequence 1040	6 AF067656	Genbank	3901271
Sequence 1040	7 AF067844	Genbank	4240386
Sequence 1040	B AF068846	Genbank	3201999
Sequence 10409	9 AF069378	Genbank	39 <b>8</b> 273 <b>3</b>
Sequence 10410	0 AF070598	Genbank	3387976
Sequence 1041		Genbank	3283905
Sequence 1041:		Genbank	3283917
Sequence 1041:		Genbank	32839 <b>23</b>
Sequence 1041		Genbank	4454685
Sequence 1041	5 AF070659	Genbank	4454693
Sequence 1041	6 AF073298	Genbank	3641537
Sequence 1041		Genbank	4689107
Sequence 1041	8 AF077036	Genbank	468911 <b>9</b>
Sequence 1041		Genbank	4689121
Sequence 1042		Genbank	4689133
Sequence 1042		≘nbank	4679017
Sequence 1042		anbank	4322311
Sequence 1042		.∋nbank	3420928
Sequence 1042		Genbank	3649 <b>656</b>
Sequence 1042		Genbank	5813822
Sequence 1042		Genbank	5114048
Sequence 1042		Genbank	5114052
Sequence 1042		Genbank	348 <b>3188</b>
Sequence 1042		Genbank	3483435
Sequence 1043		Genbank	3483552
Sequence 1043	1 AF086210	Genbank	3483555
Sequence 1043		Genbank	3483579
Sequence 1043		Genbank	3483723
Sequence 1043		Genbank	3483733
Sequence 1043		Genbank	3483902
Sequence 1043		Genbank	4322487
Sequence 1043		Genbank	4191345
Sequence 1043		Genbank	3523234
Sequence 1043		Genbank	3523242
Sequence 1044		Genbank	3859979
Sequence 1044		Genbank	4426606
		Genbank	5138909
Sequence 1044			5138919
Sequence 1044		Genbank	
Sequence 1044		Genbank	4140681
Sequence 1044	5 AF097026	Genbank	6409120

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Sequence 10446	AF097514	Genbank	4808600
Sequence 10447		Genbank	3929582
Sequence 10448		Genbank	4323586
Sequence 10449		Genbank	4324471
Sequence 10450		Genbank	4378528
		Genbank	
Sequence 10451			4139227
Sequence 10452		Genbank	4193945
Sequence 10453		Genbank	4324698
Sequence 10454		Genbank	5326796
Sequence 10455	AF112215	Genbank	6563217
Sequence 10456	AF112218	Genbank	6563223
Sequence 10457	7 AF112222	Genbank	6563 <b>229</b>
Sequence 10458	3 AF113016	Genbank	6642755
Sequence 10459		Genbank	6855 <b>63</b> 4
Sequence 10460		Genbank	6563235
Sequence 10461		Genbank	4886909
Sequence 10462		Genbank	6650827
Sequence 10463		Genbank	4903275
Sequence 10464		Genbank	4689265
Sequence 10465		Genbank	5059061
Sequence 10466		Genbank	5106995
Sequence 10467		Genbank	5106999
Sequence 10468		Genbank	6563273
Sequence 10469	AF126782	Genbank	6318547
Sequence 10470		Gen <b>ba</b> nk	6469610
Sequence 10471		Genbank	4928043
Sequence 10472		Genbank	4337095
Sequence 10473		Genbank	4558 <b>506</b>
Sequence 10474		Genbank	4406548
Sequence 10475		Genbank	4406677
Sequence 10476		Genbank	4406704
Sequence 10477		Genbank	4680 <b>646</b>
Sequence 10478		Genbank	4680 <b>648</b>
Sequence 10479	AF132956	Genbank	4680682
Sequence 10480	AF132 <b>973</b>	Genbank	468071 <b>6</b>
Sequence 10481	AF132 <b>984</b>	Genbank	4959 <b>56</b> 7
Sequence 10482		Genbank	4529886
Sequence 10483		Genbank	5006628
Sequence 10484		Genbank	5020073
Sequence 10485		Genbank	4761681
Sequence 10486		Genbank	7106777
Sequence 10487		Genbank	7106815
Sequence 10488		Genbank	6166337
Sequence 10489		Genbank	4929566
Sequence 10490		Genbank	4929616
Sequence 10491		Genbank	4929644
Sequence 10492		Genbank	4929674
Sequence 10493		Genbank	4929690
Sequence 10494		Genbank	4929726
Sequence 10495		Genbank	
			4929762
Sequence 10496		Genbank	5456956
Sequence 10497		Genbank	4929830
Sequence 10498		Genbank	5731112
Sequence 10499		Genbank	6841233
Sequence 10500		Genbank	6841255
Sequence 10501	AF161425	Genbank	68412 <b>63</b>

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<u>Table 3-5</u>

Sequence 10502	AF161428	Genbank	6841 <b>269</b>
Sequence 10503	AF161448	Genbank	6841309
Sequence 10504			
•	AF161485	Genbank	6841493
Sequence 10505	AF161490	Genbank	6841503
Sequence 10506	AF161503	Genbank	6841529
Sequence 10507	AF161556	Genbank	6841379
Sequence 10508	AF161800	Genbank	5353769
Sequence 10509	AF167570	Genbank	5762314
Sequence 10510	AF168956	Genbank	5702387
Sequence 10511	AF172368	Genbank	5679370
Sequence 10512	AF176796	Genbank	575 <b>9308</b>
Sequence 10513	AF179274	Genbank	5911402
Sequence 10514	AF180473	Genbank	6 <b>856202</b>
Sequence 10515	AF186249	Genbank	657 <b>2947</b>
Sequence 10516	AF187554	Genbank	6653225
Sequence 10517	AF190465	Genbank	612 <b>0105</b>
Sequence 10518	AF196779	Genbank	6180170
Sequence 10519	AF199339	Genbank	6708280
Sequence 10520	AF203815	Genbank	697 <b>9641</b>
Sequence 10521	AF207664	Genbank	6685071
Sequence 10522	AF213884	Genbank	7012904
Sequence 10523	AF217403	Genbank	6691160
	AJ000519		
Sequence 10524		Genbank	2739214
Sequence 10525	AJ000881	Genbank	2924308
Sequence 10526	AJ001443	Genbank	6006514
Sequence 10527	AJ001866	Genbank	29 <b>69902</b>
Sequence 10528	AJ001981	Genbank	25 <b>46978</b>
Sequence 10529	AJ002030	Genbank	2 <b>570006</b>
Sequence 10530	AJ006291	Genbank	3 <b>805937</b>
Sequence 10531	AJ010069	Genbank	34 <b>83012</b>
Sequence 10532	AJ010597	Genbank	35 <b>59873</b>
Sequence 10533	AJ011497	Genbank	4128014
Sequence 10534	AJ011930	Genbank	3859769
Sequence 10535	AJ012409	Genbank	3881975
Sequence 10536	AJ133812	Genbank	4704217
Sequence 10537	AJ223352	Genbank	325 <b>5996</b>
Sequence 10537		Genbank	
	AJ224442	T	2911586
Sequence 10539	AJ245719	Genbank	6688168
Sequence 10540	AK000023	Genbank	7019837
Sequence 10541	AK000028	Genbank	7019844
Sequence 10542	AK000086	Genbank	7019 <b>944</b>
Sequence 10543	AK000107	Genbank	7019 <b>979</b>
Sequence 10544	AK000221	Genbank	7020163
Sequence 10545	AK000291	Genbank	7020274
Sequence 10546	AK000318	Genbank	7020322
Sequence 10547	AK000348	Genbank	70 <b>20373</b>
Sequence 10548	AK000395	Genbank	7020455
Sequence 10549	AK000414	Genbank	7020486
Sequence 10550	AK000414	Genbank	70205 <b>0</b> 5
Sequence 10550		Genbank	
•	AK000434		7020520
Sequence 10552	AK000439	Genbank	7020528
Sequence 10553	AK000474	Genbank	7020586
Sequence 10554	AK000486	Genbank	7020607
Sequence 10555	AK000507	Genbank	702 <b>0644</b>
Sequence 10556	AK000541	Genbank	702 <b>0705</b>
Sequence 10557	AK000548	Genbank	7020717

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Sequence 105	58 AK0	00560	Genbank	7020738
Sequence 105		00571	Genbank	7020756
Sequence 105		00585	Genbank	7020779
Sequence 105		00667	Genbank	7020906
Sequence 105		00756	Genbank	7021041
Sequence 105		00790	Genbank	7021091
Sequence 105		00915	Genbank	7021878
Sequence 105		00934	Genbank	7021978
Sequence 105		01061	Genbank	7021909
Sequence 105		01091	Genbank	
				7022140
Sequence 105		01207	Genbank	7022315
Sequence 105		01313	Genbank	7022490
Sequence 105		01314	Genbank	7022491
Sequence 105		01321	Genbank	7022503
Sequence 105		01375	Genbank	7022596
Sequence 105		01387	Genbank	7022615
Sequence 105		01395	Genbank	7022625
Sequence 105		01413	Genbank	7022655
Sequence 105		01430	Genbank	7022682
Sequence 105		01441	Genbank	7022699
Sequence 105		01448	Genbank	7022713
Sequence 105		01451	Genbank	7022717
Sequence 105		01452	Genbank	7022718
Sequence 105		01529	Genbank	7022 <b>840</b> °
Sequence 105		01571	Genbank	7022 <b>906</b>
Sequence 105	83 AK00	01576	Genbank	7022913
Sequence 105	84 AK00	01607	Genbank	7022965
Sequence 105		01614	Genbank	7022977
Sequence 105	86 AK00	01691	Genbank	7023106
Sequence 105	87 AK00	01732	Genbank	7023176
Sequence 105		01755	Genbank	7023219
Sequence 105		01762	Genbank	7023231
Sequence 105		02036	Genbank	7023674
Sequence 1059	91 AK00	02072	Genbank	7023732
Sequence 105		8725	Genbank	2791551
Sequence 1059	93 ALO	9181	Genbank	2853179
Sequence 1059	94 AL02	21368	Genbank	3080468
Sequence 105	95 AL02	21391	Genbank	4467344
Sequence 105		21 <b>807</b>	Genbank	9367202
Sequence 105	97 AL02	21938	Genbank	3242170
Sequence 105	98 AL02	2069	Genbank	3256174
Sequence 105	99 AL02	2240	Genbank	4826471
Sequence 106	00 AL02	2312	Genbank	4914501
Sequence 106		2313	Genbank	4200326
Sequence 106		2328	Genbank	5263010
Sequence 1066		2394	Genbank	9581758
Sequence 106	04 AL02	3656	Genbank	5679746
Sequence 106		4474	Genbank	3395511
Sequence 106		4493	Genbank	3288439
Sequence 106		0996	Genbank	36 <b>88349</b>
Sequence 106		31178	Genbank	3550040
Sequence 1066		31390	Genbank	5002609
Sequence 106		31652	Genbank	9368398
Sequence 106		31659	Genbank	9 <b>581755</b>
Sequence 106		1670	Genbank	4469083
Sequence 106		1685	Genbank	9368423
Coquence 100	יט אנטט		Genbank	000420

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Sequence 10614	AL033397	Genbank	4902626
Sequence 10615		Genbank	5763714
Sequence 10616		Genbank	6807582
Sequence 10617		Genbank	4938279
Sequence 10618		Genbank	4455565
Sequence 10619		Genbank	5918013
Sequence 10620	AL034400	Genbank	4455461
Sequence 10621	AL034548	Genbank	7263904
Sequence 10622	AL034582	Genbank	5830348
Sequence 10623	AL035209	Genbank	4160217
Sequence 10624		Genbank	4200231
Sequence 10625		Genbank	6010110
Sequence 10626		Genbank	5263001
Sequence 10627		Genbank	6624641
•		Genbank	9581764
Sequence 10628			
Sequence 10629		Genbank	6015535
Sequence 10630		Genbank	4826515
Sequence 10631	AL049280	Genbank	4500037
Sequence 10632		Genbank	4500222
Sequence 10633		Genbank	4902753
Sequence 10634		Genbank	5679448
Sequence 10635	AL049712	Genbank	5629919
Sequence 10636	AL049779	Genbank	8176894
Sequence 10637		Genbank	6273553
Sequence 10638		Genbank	8217859
Sequence 10639		Genbank	6729561
Sequence 10640		Genbank	5708093
Sequence 10641	AL049843	Genbank	5830432
Sequence 10642		Genbank	4884375
		Genbank	4886440
Sequence 10643			
Sequence 10644		Genbank	4886500
Sequence 10645		Genbank	5668655
Sequence 10646		Genba	6137008
Sequence 10647		Genba	4914597
Sequence 10648		Genbar	4914585
Sequence 10649		Genbank	4914613
Sequence 10650	AL078621	Genbank	6013067
Sequence 10651	AL079295	Genbank	5102607
Sequence 10652	AL079312	Genbank	5 <b>10289</b> 0
Sequence 10653	AL079342	Genbank	6018784
Sequence 10654	AL080080	Genbank	5262491
Sequence 10655		Genbank	5262498
Sequence 10656		Genbank	5262499
Sequence 10657		Genbank	5262504
Sequence 10658		Genbank	5262506
Sequence 10659		Genbank	5262583
Sequence 10660		Genbank	
			5262652
Sequence 10661		Genbank	5870622
Sequence 10662		Genbank	57 <b>63</b> 753
Sequence 10663		Genbank	5830430
Sequence 10664		Genbank	5912599
Sequence 10665		Genbank	7228319
Sequence 10666		Genbank	61 <b>36931</b>
Sequence 10667	AL109754	Genbank	6911635
Sequence 10668	AL109758	Genbank	8217865
Sequence 10669	AL109925	Genbank	7105935
	_		

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Sequence 10670	AL110183	Genbank	5817095
Sequence 10671	AL110185	Genbank	5817098
Sequence 10672	AL110202	Genbank	5817121
Sequence 10673	AL117350	Genbank	7159817
Sequence 10674	AL117461	Genbank	5911922
Sequence 10675	AL117481	Genbank	5911958
Sequence 10676	AL117694	Genbank	8248721
Sequence 10677	AL121601	Genbank	7159760
Sequence 10678	AL121694	Genbank	6318179
Sequence 10679	AL121716	Genbank	7159797
Sequence 10680	AL122003	Genbank	6580480
Sequence 10681	AL132821	Genbank	7159786
Sequence 10682	AL132992	Genbank	6996086
Sequence 10683	AL133060	Genbank	
	AL133163		6453483
Sequence 10684		Genbank	6562367
Sequence 10685	AL133216	Genbank	6983473
Sequence 10686	AL133245	Genbank	7159621
Sequence 10687	AL133396	Genbank	6562003
Sequence 10688	AL133585	Genbank	6599171
Sequence 10689	AL135744	Genbank	6682291
Sequence 10690	AL135784	Genbank	6911343
Sequence 10691	AL135858	Genbank	7159622
Sequence 10692	AL136059	Genbank	7106618
Sequence 10693	AL136296	Genbank	8217913
Sequence 10694	AL136543	Genbank	6807646
Sequence 10695	AL136593	Genbank	7018431
Sequence 10696	AL136884	Genbank	6 <b>807672</b>
Sequence 10697	AL137229	Genbank	70 <b>09597</b>
Sequence 10698	AL137440	Genbank	6 <b>808001</b>
Sequence 10699	AL137489	Genbank	6808110
Sequence 10700	AL137496	Genbank	6808137
Sequence 10701	AL137519	Genbank	6 <b>808181</b>
Sequence 10702	AL137543	Genbank	6 <b>808222</b>
Sequence 10703	AL137686	Genbank	6 <b>807956</b>
Sequence 10704	AL137692	Genbank	6 <b>808008</b>
Sequence 10705	AL137751	Genbank	68 <b>08387</b>
Sequence 10706	AL137818	Genbank	7009598
Sequence 10707	AP000046	Genbank	3132356
Sequence 10708	AP000085	Genbank	4730826
Sequence 10709	AP000217	Genbank	4827264
Sequence 10710	AP000345	Genbank	51 <b>03008</b>
Sequence 10711	AP000350	Genbank	5103013
Sequence 10712	AP000352	Genbank	6016843
Sequence 10713	AP000500	Genbank	5 <b>926687</b>
Sequence 10714	AP000525	Genbank	5931 <b>503</b>
Sequence 10715	AP000700	Genbank	6705908
Sequence 10716	AP000962	Genbank	6942330
Sequence 10717	AP001052	Genbank	6693602
Sequence 10718	AP001136	Genbank	697 <b>0360</b>
Sequence 10719	AP001137	Genbank	69 <b>70361</b>
Sequence 10720	AP001138	Genbank	7262569
Sequence 10721	AP001172	Genbank	6983884
Sequence 10722	AP001252	Genbank	7077183
Sequence 10723	AP001343	Genbank	7209831
Sequence 10724	AP001346	Genbank	7209834
Sequence 10725	D13866	Genbank	433410

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Sequence 10726	D13900	Genbank	433412
Sequence 10727	D14043	Genbank	219924
Sequence 10728	D14658	Genbank	285940
Sequence 10729	D14662	Genbank	285948
Sequence 10730	D14665	Genbank	6630617
Sequence 10731	D14696	Genbank	285962
Sequence 10732	D14710	Genbank	559324
Sequence 10733	D15057	Genbank	493244
Sequence 10734	D16111	Genbank	435637
Sequence 10735	D16234	Genbank	
	D16481	Genbank	303617
Sequence 10736			473711
Sequence 10737	D16938	Genbank	598868
Sequence 10738	D16947	Genbank	598932
Sequence 10739	D21089	Genbank	475156
Sequence 10740	D25218	Genbank	434778
Sequence 10741	D26600	Genbank	565650
Sequence 10742	D29805	Genbank	474986
Sequence 10743	D38112	Genbank	644480
Sequence 10744	D42041	Genbank	577294
Sequence 10745	D42052	Genbank	1136740
Sequence 10746	D42063	Genbank	924266
Sequence 10747	D45915	Genbank	1483130
Sequence 10748	D49355	Genbank	1020405
Sequence 10749	D50372	Genbank	2605593
Sequence 10750	D50929	Genbank	1469200
Sequence 10751	D63486	Genbank	1469885
Sequence 10752	D63874	Genbank	968887
Sequence 10753	D66904	Genbank	1235726
Sequence 10754	D67025	Genbank	2656091
Sequence 10755	D79986	Genbank	1136389
Sequence 10756	D80000	Genbank	1136415
Sequence 10757	D80007	Genbank	1136429
Sequence 10758	D83253	Genbank	2951772
Sequence 10759	D84907	Genbank	1596053
Sequence 10760	D85181	Genbank	1906795
Sequence 10761	D86962	Genbank	1503997
Sequence 10762	D87018	Genbank	2114280
Sequence 10763	D87127	Genbank	1817551
Sequence 10764	D87666	Genbank	1620016
Sequence 10765	D87742	Genbank	1665824
Sequence 10766	D89052	Genbank	1694672
Sequence 10767	E01716	Genbank	2169969
Sequence 10768	E01932	Genbank	2170180
Sequence 10769	E01954	Genbank	2170100
Sequence 10709	E02628		
Sequence 10770	E13330	Genbank Genbank	2170856
Sequence 10771	J01415		3252135
		Genbank	1944628
Sequence 10773	J03077	Genbank	183230
Sequence 10774	J03473	Genbank	337423
Sequence 10775	J03746	Genbank	183655
Sequence 10776	J04799	Genbank	190197
Sequence 10777	J04823	Genbank	1311703
Sequence 10778	J05016	Genbank	181507
Sequence 10779	J05480	Genbank	179807
Sequence 10780	L01057	Genbank	292462
Sequence 10781	L04483	Genbank	292444

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Sequence 10	0782	L05092	Genbank	388031
Sequence 10	0783	L05628	Genbank	1835658
Sequence 10		L08125	Genbank	306500
Sequence 10		L09159	Genbank	307374
Sequence 10		L11370	Genbank	387674
•			Genbank	
Sequence 10		L12136		181536
Sequence 10		L12168	Genbark	178083
Sequence 10		L19185	Genbark	440307
Sequence 10	0790	L20941	Genbank	507251
Sequence 10		L28809	Genbank	454151
Sequence 10	0792	L29252	Genbank	808011
Sequence 10	0793	L31 <b>61</b> 0	Genbank	1220360
Sequence 10		L34160	Genbank	642091
Sequence 10		L35681	Genbank	532031
Sequence 10		L38933	Genbank	1008841
Sequence 10		L38951	Genbank	893287
Sequence 10		L38995	Genbank	704415
•		L39068	Genbank	994714
Sequence 10				
Sequence 10		L39945	Genbank	703081
Sequence 10		L40407	Genbank	703117
Sequence 10		L41887	Genbank	950423
Sequence 10		L47647	Genbank	1000861
Sequence 10	0804	L48984	Genbank	1066728
Sequence 10	0805	L78754	Genbank	1374832
Sequence 10	0806	M11147	Genbank	182513
Sequence 10	0807	M15887	Genbank	181960
Sequence 10		M16342	Genbank	184266
Sequence 10		M17886	Genbank	190233
Sequence 10		M18930	Genbank	184371
Sequence 10		M19989	Genbank	189719
Sequence 10		M19997	Genbank	181968
Sequence 10		M20496	Genbank	809235
Sequence 10		M21142	Genbank	183402
Sequence 10		M21154	Genbank	178517
Sequence 10		M21575	Genbank	1311702
Sequence 10		M24543	Genbank	341200
				189618
Sequence 10		M24902	Genbank	
Sequence 10		M25639	Genbank	188555
Sequence 10		M26481	Genbank	619789
Sequence 10		M26663	Genbank	618463
Sequence 10		M27024	Genbank	341598
Sequence 10		M29064	Genbank	337452
Sequence 10	0824	M29366	Genbank	181979
Sequence 10	0825	M29877	Genbank	178408
Sequence 10	0826	M31520	Genbank	337504
Sequence 10	0827	M33146	Genbank	181070
Sequence 10	0828	M33616	Genbank	184381
Sequence 10		M34600	Genbank	180934
Sequence 10		M36341	Genbank	178984
Sequence 10		M37583	Genbank	184059
Sequence 10		M58459	Genbank	337511
Sequence 10	1833	M58485	Genbank	180154
Sequence 10		M60457	Genbank	181249
Sequence 10		M60756	Genbank	184085
Sequence 10		M64929	Genbank	190421
Sequence 10	J <b>53</b> /	M59181	Genbank	641957

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Sequence 10838	M74775	Genbank	187151
Sequence 10839	M76299	Genbank	188778
Sequence 10840	M81182	Genbank	190128
Sequence 10841	M81601	Genbank	339442
Sequence 10842	M83088	Genbank	189925
Sequence 10843			
	M93036	Genbank	182904
Sequence 10844	M95775	Genbank	187180
Sequence 10845	M96256	Genbank	182625
Sequence 10846	M96982	Genbank	338262
Sequence 10847	S48220	Genbank	257451
Sequence 10848	S67861	Genbank	239579
Sequence 10849	S70290	Genbank	546602
Sequence 10850	S72481	Genbank	632789
Sequence 10851	S75463	Genbank	833998
Sequence 10852	U00968	Genbank	409404
Sequence 10853	U01925	Genbank	460086
Sequence 10854	U09582	Genbank	493079
Sequence 10855	U09646	Genbank	1041191
Sequence 10856	U09813	Genbank	
•	U12404		1008454
Sequence 10857		Genbank	531170
Sequence 10858	U12465	Genbank	562073
Sequence 10859	U14650	Genbank	541612
Sequence 10860	U14970	Genbank	550020
Sequence 10861	U15008	Genbank	600747
Sequence 10862	U16797	Genbank	902370
Sequence 10863	U17899	Genbank	717053
Sequence 10864	U19144	Genbank	914902
Sequence 10865	U21090	Genbank	1008457
Sequence 10866	U23803	Genbank	773643
Sequence 10867	U25182	Genbank	799380
Sequence 10868	U28369	Genbank	974283
Sequence 10869	U30826	Genbank	1049079
Sequence 10870	U32944	Genbank	1209060
Sequence 10871	U <b>36336</b>	Genbank	1209628
Sequence 10872	U38178	Genbank	1145301
Sequence 10873	U44754	Genbank	1174202
Sequence 10874	U47742	Genbank	15 <b>179</b> 13
Sequence 10875	U49245	Genbank	1216503
Sequence 10876	U49973	Genbank	2226003
Sequence 10877	U50939	Genbank	1314559
Sequence 10878	U52111	Genbank	8331754
Sequence 10879	U54993	Genbank	4097314
Sequence 10880	U55937	Genbank	1575044
Sequence 10881	U57847	Genbank	1373420
Sequence 10882	U58522	Genbank	1381163
Sequence 10883	U60644	Genbank	1575346
Sequence 10884	U60808	Genbank	1915971
Sequence 10885	U62961	Genbank	1519051
•	U65896		1763689
Sequence 10886	U66871	Genbank	
Sequence 10887		Genbank	1519518
Sequence 10888	U68105	Genbank	1562509
Sequence 10889	U <b>73824</b>	Genbank	1857236
Sequence 10890	U77643	Genbank	2062390
Sequence 10891	U79273	Genbank	1710239
Sequence 10892	U79274	Genbank	1710240
Sequence 10893	U80017	Genbank	1737211

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Secuence 10894	U <b>83463</b>	Genbank	1916849
Secuence 10895	U87460	Genbank	2076881
Sequence 10896	U <b>90441</b>	Genbank	2439984
Secuence 10897	<b>⊍90915</b>	Genbank	1913895
Sequence 10898	U91323	Genbank	3582311
Sequence 10899	U91328	Genbank	2088550
Sequence 10900	U97670	Genbank	3264860
Sequence 10901	V00478	Genbank	28244
Sequence 10902	X00351	Genbank	28251
Sequence 10903	X03083	Genbank	34309
Sequence 10904	X04588	Genbank	37423
Sequence 10905	X05332	Genbank	35740
Sequence 10906	X13238	Genbank	1200056
Sequence 10907	X15187	Genbank	37260
Sequence 10907 Sequence 10908	X15341		
Sequence 10908		Genbank	1197215
	X15822	Genbank	30146
Sequence 10910	X51525	Genbank	30170
Sequence 10911	X53280	Genbank	29504
Sequence 10912	X53586	Genbank	33943
Sequence 10913	X56932	Genbank	23690
Sequence 10914	X58288	Genbank	32455
Sequence 10915	X60221	Genbank	509290
Sequence 10916	X63432	Genbank	28335
Sequence 10917	X63657	Genbank	296185
Sequence 10918	X64707	Genbank	29382
Sequence 10919	×65 <b>923</b>	Genbank	ປ1 <b>302</b>
Sequence 10920	X67247	Genbank	36149
Sequence 10921	X67951	Genbank	287640
Sequence 10922	X74801	Genbank	6715 <b>26</b>
Sequence 10923	X74968	Genbank	439659
Sequence 10924	X75 <b>821</b>	Genbank	5 <b>09780</b>
Sequence 10925	X75861	Genbank	456258
Sequence 10926	X78136	Genbank	460772
Sequence 10927	X78137	Genbank	460770
Sequence 10928	X78669	Genbank	469 <b>884</b>
Sequence 10929	X80199	Genbank	2385366
Sequence 10930	X81109	Genbank	53 <b>5057</b>
Sequence 10931	X84 <b>694</b>	Genbank	872117
Sequence 10932	X84 <b>908</b>	Genbank	1502344
Sequence 10933	X86779	Genbank	1006658
Sequence 10934	X87176	Genbank	1050516
Sequence 10935	X87949	Genbank	1143491
Sequence 10936	X91790	Genbank	2511667
Sequence 10937	X97 <b>544</b>	Genbank	1770563
Sequence 10938	X98356	Genbank	1648 <b>869</b>
Sequence 10939	X995 <b>85</b>	Genbank	1770518
Sequence 10940	Y00052	Genbank	30308
Sequence 10941	Y00282	Genbank	3 <b>6048</b>
Sequence 10942	Y00345	Genbank	355 <b>69</b>
Sequence 10943	Y00815	Genbank	34 <b>266</b>
Sequence 10944	Y13620	Genbank	2181877
Sequence 10945	Y15286	Genbank	2584788
Sequence 10946	Z11692	Genbank	31107
Sequence 10947	Z14000	Genbank	296 <b>063</b>
Sequence 10948	Z24724	Genbank	505 <b>034</b>
Sequence 10949	Z36832	Genbank	533946
		Genbank	2003-40

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<u>Table 3-5</u>

Sequence 10950	Z37986	Genbank	780262
Sequence 10951	Z37994	Genbank	556810
Sequence 10952	Z49250	Genbank	807889
Sequence 10953	Z63138	Genbank	1035516
Sequence 10954	Z64102	Genbank	1036480
Sequence 10955	Z68284	Genbank	1130698
Sequence 10956	Z70702	Genbank	1332492
Sequence 10957	Z82194	Genbank	2582744
Sequence 10958	Z83745	Genbank	1754650
Sequence 10959	Z83844	Genbank	4467204
	Z83856	Genbank	2995261
Sequence 10960	Z84722	Genbank	1817579
Sequence 10961		Genbank	2326514
Sequence 10962	Z85999		1869775
Sequence 10963	Z92544	Genbank	
Sequence 10964	Z93016	Genbank	9650703
Sequence 10965	Z93242	Genbank	3164071
Sequence 10966	Z95114	Genbank	5101742
Sequence 10967	Z95704	Genbank	2121307
Sequence 10968	Z97053	Genbank	9650676
Sequence 10969	Z97056	Genbank	2832593
Sequence 10970	Z97183	Genbank	2961 <b>406</b>
Sequence 10971	Z97630	Genbank	4582128
Sequence 10972	Z98749	Genbank	4775 <b>611</b>
Sequence 10973	Z98882	Genbank	4490865
Sequence 10974	Z98884	Genbank	5 <b>304861</b>
Sequence 10975	AA001709	EST	1445523
Sequence 10976	AA004303	EST	1447948
Sequence 10977	AA007269	EST	1463303
Sequence 10978	AA010173	EST	1471349
Sequence 10979	AA010233 ·	EST	1471260
Sequence 10980	AA010519	EST	1471545
Sequence 10981	AA011539	EST	1472565
Sequence 10982	AA015585	EST	1476651
Sequence 10983	AA017170	EST	1479335
Sequence 10984	AA018892	EST	1482283
Sequence 10985	AA021623	EST	1485284
Sequence 10986	AA025361	EST	1489375
Sequence 10987	AA026455	EST	1492355
Sequence 10988	AA026758	EST	1492556
Sequence 10989	AA027906	EST	1493993
Sequence 10990	AA031317	EST	1501272
Sequence 10991	AA035377	EST	1507101
Sequence 10992	AA037143	EST	1512251
Sequence 10993	AA039734	EST	1516236
Sequence 10994	AA039957	EST	1516261
Sequence 10995	AA040714	EST	1516992
Sequence 10996	AA044237	EST	1522113
Sequence 10997	AA044575	EST	1522808
Sequence 10998	AA044703	EST	1522971
Sequence 10999	AA045239	EST	1523443
Sequence 11000	AA045616	EST	1525379
Sequence 11001	AA047046	EST	1524944
Sequence 11002	AA054272	EST	1545197
Sequence 11003	AA057541	EST	1550246
Sequence 11004	AA057541 AA058950	EST	1551758
Sequence 11005	AA056950 AA062994	EST	1557655
Sequence 11005	MAUDZ994	EOI	1337033

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Secuence 11006	AA063190	EST.	155 <b>6924</b>
Sequence 11007	AA063198	EST	1556796
Sequence 11008	AA070463	EST	1577823
Sequence 11009	AA070785	EST	1578347
Sequence 11010	AA074614	EST	1614492
Sequence 11011	AA075478	EST	1615543
Sequence 11012	AA076016	EST	
Sequence 11012	AA083482	EST	1615903
Sequence 11013	AA085007		1625543
Sequence 11014 Sequence 11015		EST	1627125
	AA085810	EST	1629342
Sequence 11016	AA088344	EST	1633 <b>856</b>
Sequence 11017	AA089630	EST	1636 <b>122</b>
Secuence 11018	AA093542	EST	1639127
Sequence 11019	AA095108	EST	1640 <b>693</b>
Sequence 11020	AA098799	EST	1648 <b>802</b>
Sequence 11021	AA099464	EST	1645 <b>408</b>
Sequence 11022	AA099750	EST	1645 <b>86</b> 1
Sequence 11023	AA100034	EST	1647 <b>326</b>
Sequence 11024	AA101634	EST	1 <b>648570</b>
Sequence 11025	AA101663	EST	1648744
Sequence 11026	AA102383	EST	1647229
Sequence 11027	AA102507	EST	1647621
Sequence 11028	AA113782	EST	1 <b>667703</b>
Sequence 11029	AA114830	EST	1669952
Sequence 11030	AA115175	EST	1670372
Sequence 11031	AA115455	EST	1670257
Sequence 11032	AA115636	EST	1670771
Sequence 11033	AA115738	EST	1670751
Sequence 11034	AA122059	EST	167807 <b>8</b>
Sequence 11035	AA122097	EST	1678318
Sequence 11036	AA126927	EST	1687948
Sequence 11037	AA131505	EST	1692993
Sequence 11038	AA131583	EST	1693072
Sequence 11039	AA133311	EST	1690351
Sequence 11040	AA135694	EST	1696705
Sequence 11041	AA136084	EST	1697294
Sequence 11042	AA136337	EST	1697545
Sequence 11043	AA137210	EST	1697007
Sequence 11044	AA143393	EST	1712764
Sequence 11045	AA146684	EST	1716058
Sequence 11046	AA149579	EST	1720380
Sequence 11047	AA150361	EST	1721873
Sequence 11048	AA151438	EST	1719775
Sequence 11049	AA151565	EST	1720183
Sequence 11050	AA151816	EST	1720511
Sequence 11051	AA152016	EST	1720854
Sequence 11052	AA156951	EST	1728584
Sequence 11053	AA157731	EST	1729356
Sequence 11054	AA 160577	EST	1735945
Sequence 11055	AA161189	EST	1735945
Sequence 11056	AA161337		
Sequence 11057	AA165446	EST	1735652
•		EST	1741462
Sequence 11058	AA169597	EST	1747985
Sequence 11059	AA169657	EST	1748026
Sequence 11060	AA172193	EST	1751270
Sequence 11061	AA173705	EST	1753 <b>883</b>

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Sequence	11062	AA174161	EST	1754441
Secuence	11063	AA176485	EST	1757618
Sequence		AA177128	EST	1758286
Sequence		AA178986	EST	1760538
Sequence :		AA180264	EST	1761531
Sequence		AA186328	EST	1774446
Sequence		AA187076	EST	1775245
Sequence		AA188297	EST	<b>17</b> 75383
Sequence		AA190560	EST	<b>17</b> 79535
Sequence:	11071	AA193327	EST	<b>1782</b> 738
Sequence 1	11072	AA193455	EST	<b>178284</b> 6
Sequence '	11073	AA203319	EST	1799028
Sequence 1	11074	AA206446	EST	1801826
Sequence :		AA207077	EST	1802642
Sequence :		AA209329	EST	1807273
Sequence		AA211900	EST	1810537
Sequence 1		AA223613	EST	1844155
Sequence 1		AA225958	EST	1847285
Sequence :		AA226672	EST	1847988
Sequence :		AA232991	EST	1856115
Sequence		AA232991 AA233294		
Sequence	11002		EST	1856287
Sequence 1		AA234637	EST	1859306
Sequence 1		AA235298	EST	1859735
Sequence 1		AA243646	EST	1874456
Sequence 1		AA248470	EST	1879307
Sequence 1		AA248684	EST	1879900
Sequence 1		AA250936	EST	1886035
Sequence 1		AA251061	EST	1886024
Sequence 1		AA251263	EST	1886226
Sequence 1		AA251692	EST	1886655
Sequence 1		AA252143	EST	<b>188</b> 7106
Sequence 1		AA252608	EST	1887582
Sequence 1	11094	AA252943	EST	1882696
Sequence 1	11095	AA255735	EST	1892673
Sequence 1	11096	AA257016	EST	1891145
Sequence 1	11097	AA258732	EST	1893893
Sequence 1	11098	AA262091	EST	1898235
Sequence 1	11099	AA262425	EST	1897921
Sequence 1	11100	AA262683	EST	1898095
Sequence 1		AA278796	EST	1920117
Sequence 1		AA282285	EST	1925201
Sequence 1		AA285126	EST	1928107
Sequence 1		AA286765	EST	1933628
Sequence 1		AA293822	EST	1941745
Sequence 1		AA296237	EST	1948581
Sequence 1		AA296690	EST	1949182
Sequence 1		AA296812	EST	1949146
Sequence 1		AA296923	EST	1949255
Sequence 1		AA304244	EST	1956597
		AA305213		
Sequence 1		<del>-</del>	EST	1957540
Sequence 1		AA305921	EST	1958412
Sequence 1		AA306323	EST	1958671
Sequence 1		AA306941	EST	1959271
Sequence 1		AA307728	EST	1960056
Sequence 1		AA308688	EST	1961018
Sequence 1	11117	AA309119	EST	1961444

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Sequence 11118	AA309929	EST	196 <b>2484</b>
Sequence 11119	AA310375	EST	1962704
Sequence 11120	AA311577	EST	1963980
Sequence 11121	AA313294	EST	1965623
Sequence 11121	AA314382	EST	1966731
Sequence 11122			1967114
Sequence 11123	AA314635	EST	
Sequence 11124	AA315157	EST	1967486
Sequence 11125	AA315989	EST	1968318
Sequence 11126	AA316456	EST	1968785
Sequence 11127	AA316676	EST	1969004
Sequence 11128	AA325081	EST	197757 <b>9</b>
Sequence 11129	AA325713	EST	1977956
Sequence 11130	AA329695	EST	1981937
Sequence 11131	AA331249	EST	1983730
Sequence 11132	AA340295	EST	1992760
Sequence 11133	AA340744	EST	1992983
Sequence 11134	AA343834	EST	1996133
	AA345527	EST	1997837
Sequence 11135		EST	2007708
Sequence 11136	AA355326		2016963
Sequence 11137	AA364646	EST	2017902
Sequence 11138	AA365552	EST	
Sequence 11139	AA371377	EST	2023695
Sequence 11140	AA379942	EST	2032282
Sequence 11141	AA382161	EST	2034479
Sequence 11142	AA384806	EST	203 <b>7123</b>
Sequence 11143	AA393525	EST	2046493
Sequence 11144	AA400534	EST	2054467
Sequence 11145	AA400644	EST	2054515
Sequence 11146	AA402038	EST	2056038
Sequence 11147	AA402743	EST	2057061
Sequence 11148	AA402800	EST	2056553
Sequence 11149	AA404235	EST	2058977
Sequence 11150	AA404535	EST	2059268
Sequence 11151	AA406310	EST	2064294
Sequence 11152	AA410936	EST	2070190
Sequence 11153	AA411736	EST	2069397
Sequence 11155	AA417324	EST	2077423
Sequence 11154	AA417743	EST	2079544
Sequence 11155	AA418847	EST	2080666
Sequence 11156			2094559
Sequence 11157	AA420643	EST	2094538
Sequence 11158	AA420650	EST	2102976
Sequence 11159	AA424006	EST	
Sequence 11160	AA424349	EST	2103336
Sequence 11161	AA425535	EST	2106293
Sequence 11162	AA426021	EST	2106545
Sequence 11163	AA427560	EST	2111428
Sequence 11164	AA428565	EST	2112562
Sequence 11165	AA430605	EST	21 <b>11178</b>
Sequence 11166	AA432282	EST	2114670
Sequence 11167	AA436100	EST	2141014
Sequence 11168	AA437224	EST	2142138
Sequence 11169	AA442070	EST	2153948
Sequence 11170	AA443297	EST	2155972
Sequence 11171	AA443762	EST	2156437
Sequence 11172	AA446702	EST	2159367
Sequence 11173	AA446759	EST	2159424
Sequence 111/3	W-440108	231	L 1007L4

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Sequence 11174	AA446859	EST	2159524
Sequence 11175	AA446942	EST	2159607
Sequence 11176	AA447406	EST	2161076
Sequence 11177	AA447871	EST	2161541
Sequence 11178	AA448125	EST	2161795
Sequence 11179	AA448292	EST	2161962
Sequence 11180	AA449087	EST	2163107
Sequence 11181	AA450129	EST	2163879
Sequence 11182	AA451779	EST	2165448
Sequence 11183	AA452120	EST	2165789
Sequence 11184	AA454989	EST	2177765
Sequence 11185	AA455869	EST	2178645
Sequence 11186	AA456454	EST	2179030
Sequence 11187	AA460766	EST	2185886
Sequence 11188	AA464281	EST	2189165
Sequence 11189	AA464409	EST	2189293,
Sequence 11190	AA464803	EST	2189687
Sequence 11191	AA468360	EST	2194894
Sequence 11192	AA476267	EST	2204478
Sequence 11193	AA478669	EST	2207303
Sequence 11194	AA483131	EST	2211380
Sequence 11195	AA483251	EST	2212064
Sequence 11196	AA483450	EST	2212263
Sequence 11197	AA484155	EST	2212968
Sequence 11198	AA484907	EST	2213974
Sequence 11199	AA485065	EST	2214284
Sequence 11200	AA485608	EST	2214827
Sequence 11201	AA487694	EST	2217858
Sequence 11202	AA489068	EST	2218670
Sequence 11203	AA489782	EST	2220666
Sequence 11204	AA491858	EST	2221420
Sequence 11205	AA492143	EST	2221705
equence 11206	AA492423	EST	2221985
Sequence 11207	AA493522	EST	2223363
Sequence 11208 Sequence 11209	AA493635	EST	2223476
Sequence 11210	AA502361	EST	2237328
Sequence 11211	AA502517 AA503033	EST	2237484
Sequence 11212	AA503033	EST	2238000
Sequence 11213	AA504572	EST EST	2238045
Sequence 11214	AA504890	EST	2240732
Sequence 11215	AA508394	EST	2241050 2245897
Sequence 11216	AA508597	EST	
Sequence 11217	AA513123	EST	2246100 2251535
Sequence 11218	AA516472	EST	2255996
Sequence 11219	AA523172	EST	2263884
Sequence 11220	AA524187	EST	2265115
Sequence 11221	AA525960	EST	2268029
Sequence 11222	AA526740	EST	2268809
Sequence 11223	AA527181	EST	2269250
Sequence 11224	AA527494	EST	2269563
Sequence 11225	AA527667	EST	2269736
Sequence 11226	AA528148	EST	2270217
Sequence 11227	AA531585	EST	2274291
Sequence 11228	AA532665	EST	2276919
Sequence 11229	AA532805	EST	2278381

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Secuence 11230         AA533279         EST         2277375           Secuence 11231         AA535098         EST         2279351           Secuence 11233         AA551347         EST         2287971           Secuence 11234         AA551385         EST         2321933           Secuence 11236         AA551709         EST         2321961           Secuence 11237         AA551770         EST         2322022           Secuence 11238         AA557174         EST         2327651           Sequence 11239         AA557174         EST         2327661           Sequence 11240         AA559906         EST         2331397           Secuence 11241         AA559906         EST         2331397           Secuence 11241         AA563778         EST         2335767           Sequence 11242         AA564118         EST         2335767           Sequence 11243         AA565432         EST         2334826           Sequence 11244         AA569432         EST         2348236           Sequence 11245         AA579486         EST         2357670           Sequence 11246         AA579486         EST         2359612           Sequence 11248         AA582006         <					
Secuence 11231         AA535098         EST         2279351           Secuence 11232         AA541537         EST         2287971           Secuence 11234         AA551141         EST         2321333           Secuence 11235         AA551709         EST         2321637           Secuence 11236         AA551770         EST         2327651           Sequence 11237         AA557174         EST         2327651           Sequence 11238         AA557191         EST         2327651           Sequence 11240         AA559906         EST         2331397           Sequence 11240         AA559906         EST         2331397           Sequence 11241         AA563778         EST         2335117           Sequence 11243         AA565432         EST         2337071           Sequence 11244         AA569846         EST         2337071           Sequence 11244         AA569846         EST         2348236           Sequence 11244         AA569846         EST         2357670           Sequence 11247         AA580489         EST         2356684           Sequence 11247         AA580489         EST         2359612           Sequence 11250         AA582370         <	Secuence 1	11230	AA533279	EST	2277375
Secuence 11232         AA541537         EST         2287971           Secuence 11233         AA551141         EST         2321393           Secuence 11236         AA551709         EST         2321637           Secuence 11236         AA551770         EST         2322022           Secuence 11238         AA551774         EST         2322651           Sequence 11239         AA557191         EST         2327668           Sequence 11240         AA5597633         EST         2331397           Sequence 11241         AA559906         EST         2331397           Sequence 11241         AA569178         EST         2335757           Sequence 11243         AA565432         EST         2337071           Sequence 11244         AA569846         EST         2337677           Sequence 11245         AA579486         EST         23357670           Sequence 11246         AA579486         EST         2356684           Sequence 11247         AA580489         EST         2356684           Sequence 11249         AA582252         EST         2356684           Sequence 11251         AA583370         EST         2356612           Sequence 11251         AA583922					
Secuence 11233         AA551141         EST         2321393           Secuence 11234         AA551385         EST         2321637           Secuence 11236         AA551779         EST         2322022           Secuence 11237         AA557174         EST         2327651           Sequence 11239         AA557191         EST         2327651           Sequence 11240         AA559906         EST         2331397           Sequence 11241         AA569906         EST         2335757           Sequence 11242         AA56432         EST         2335757           Sequence 11243         AA565432         EST         2337071           Sequence 11244         AA569846         EST         2337071           Sequence 11244         AA569846         EST         2337670           Sequence 11244         AA569846         EST         2357670           Sequence 11247         AA580489         EST         2357670           Sequence 11247         AA580489         EST         2355816           Sequence 11247         AA582006         EST         2359730           Sequence 11250         AA582370         EST         2359612           Sequence 11250         AA589402 <t< td=""><td></td><td></td><td></td><td></td><td></td></t<>					
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**Table 3-5** 

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<u>Table 3-5</u>

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<u>Table 3-5</u>

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Sequence 11480	Al359812	EST	4111433
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Sequence 11482	Al366222	EST	4125911
Sequence 11483	Al393262	EST	4222 <b>809</b>
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Sequence 11486	AI417686	EST	4261190
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Sequence 11488	Al436691	EST	4283721
Sequence 11489	Al439350	EST	4303707
Sequence 11490	AI468004	EST	4330094
Sequence 11491	Al469347	EST	4331437
Sequence 11492	Al469505	EST	4331595
Sequence 11493	Al469600	EST	4331690
Sequence 11494	Al472201	EST	4334291
Sequence 11495	AI47B202	EST	4371428
Sequence 11496	AI479751	EST	4372919
Sequence 11497	Al480256	EST	4373424
Sequence 11498	Al497917	EST	4389899
Sequence 11499	Al499665	EST	4391647
Sequence 11500	AI521246	EST	4435381
Sequence 11501	AI525607	EST	4439742
Sequence 11502	AI539780	EST	4453915
Sequence 11502	Al559444	EST	4509649
Sequence 11504	AI559444 AI559531		
		EST	450 <b>9736</b>
Sequence 11505	A1565555	EST	4524012
Sequence 11506	AI570526	EST	4533900
Sequence 11507	AI582754	EST	456 <b>865</b> 1
Sequence 11508	A1608932	EST	4618099
Sequence 11509	AI608936	EST	4618 <b>103</b>

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<u>Table 3-5</u>

Sequence 1151	O Al621173	EST	4630299
Sequence 1151		EST	4648794
Sequence 1151		EST	4650083
Sequence 1151		EST	4650750
Sequence 1151		EST	4683326
Sequence 1151		EST	4684285
Sequence 1151		EST	4686727
Sequence 1151		EST	
Sequence 1151		EST	4687086
Sequence 1151			4741090
Sequence 1151		EST	4893981
		EST	4896054
Sequence 1152		EST	4987014
Sequence 1152		EST	4988556
Sequence 1152		EST	5056362
Sequence 1152		EST	5109477
Sequence 1152		EST	5112262
Sequence 1152		EST	5112630
Sequence 1152		EST	5129373
Sequence 1152		EST	5129439
Sequence 1152		EST	5178162
Sequence 1153		EST	5179 <b>011</b>
Sequence 1153		EST	5232199
Sequence 1153:		EST	5365467
Sequence 1153:	3 AI802542	EST	5368014
Sequence 1153	4 Al804082	EST	5369554
Sequence 1153	5 AI808503	EST	5395069
Sequence 1153	6 Al813654	EST	5424869
Sequence 1153	7 Al815499	EST	5431045
Sequence 1153	B AI8580 <b>89</b>	EST	5511705
Sequence 11539	9 AI866603	EST	5530710
Sequence 11540	O AI866803	EST	5530834
Sequence 1154	1 AI8674 <b>6</b> 4	EST	5540480
Sequence 1154	2 AI879010	EST	5553059
Sequence 11543	3 AI887361	EST	5592525
Sequence 1154	4 AI903990	EST	6494286
Sequence 11549	5 AI905002	EST	6495389
Sequence 1154		EST	5665321
Sequence 11547	7 Al9294 <b>83</b>	EST	5665447
Sequence 11548	3 Al948496	EST	5740806
Sequence 11549	Al961342	EST	5754055
Sequence 11550	AI983520	EST	5810739
Sequence 1155		EST	4727122
Sequence 1155		EST	5405665
Sequence 1155		EST	5405971
Sequence 11554	4 AL037828	EST	5407161
Sequence 11558		EST	5408788
Sequence 11556		EST	5422386
Sequence 11557		EST	5432220
Sequence 11558		EST	4727722
Sequence 11559		EST	5925144
Sequence 11560		EST	5926825
Sequence 1156		EST	6603054
Sequence 11562		EST	6603234
Sequence 11563		EST	6854954
Sequence 11564		EST	6855132
Sequence 11565		EST	5850564
Soquence 11000	/111000540		3030304

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Sequence 11566	AW008029	EST	5856807
Sequence 11567	AW008048	EST	
Sequence 11568	AW008782		5856826
Sequence 11569		EST	5857560
	AW021322	EST	5874782
Sequence 11570	AW021588	EST	5875118
Sequence 11571	AW022159	EST	5875689
Sequence 11572	AW043631	EST	5904160
Sequence 11573	AW044310	EST	5904839
Sequence 11574	AW069285	EST	6024283
Sequence 11575	AW069315	EST	6024313
Sequence 11576	AW073580	EST	6028578
Sequence 11577	AW082238	EST	6037390
Sequence 11578	AW083099	EST	6038251
Sequence 11579	AW085581	EST	6040733
Sequence 11580	AW087200	EST	6043005
Sequence 11581	AW136642	EST	6140775
Sequence 11582	AW148690	EST	6196586
Sequence 11583	AW157858	EST	6229259
Sequence 11584	AW160706		
Sequence 11585	AW161328	EST	6299739
Sequence 11586	AW 161326 AW 162943	EST EST	6300361
			6301976
Sequence 11587	AW167129	EST	6398654
Sequence 11588	AW168050	EST	6399575
Secuence 11589	AW168184	EST	6399709
Sequence 11590	AW168400	EST	6399925
Sequence 11591	AW177050	EST	6443087
Sequence 11592	AW192594	EST	6471293
Sequence 11593	AW205906	EST	6505380
Sequence 11594	AW247236	EST	6590229
Sequence 11595	AW250104	EST	6593 <b>097</b>
Sequence 11596	AW294606	EST	6701242
Sequence 11597	AW294626	EST	67 <b>01262</b>
Sequence 11598	AW297943	EST	6704579
Sequence 11599	AW302841	EST	6712521
Sequence 11600	AW337419	EST	6834045
Sequence 11601	AW363949	EST	6868599
Sequence 11602	AW 363975	EST	6868625
Sequence 11603	AW365134	EST	68 <b>69784</b>
Sequence 11604	AW365960	EST	6870610
Sequence 11605	AW368992	EST	6873642
Sequence 11606	AW376614	EST	6881275
Sequence 11607	AW380617	EST	688 <b>5276</b>
Sequence 11608	AW380631	EST	68 <b>85290</b>
Sequence 11609	AW381959	EST	688 <b>6618</b>
Sequence 11610	AW386920	EST	6891579
Sequence 11611	AW401953	EST	6920639
Sequence 11612	AW402499	EST	6921188
Sequence 11613	AW450642	EST	6991418
Sequence 11614	AW452921	EST	6993 <b>69</b> 7
Sequence 11615	AW499890	EST	7111988
Sequence 11616	AW503119	EST	7118224
Sequence 11617	AW518064	EST	N/A
Sequence 11618	AW 573593	EST	7238326
Sequence 11619	AW574740	EST	7246279
Sequence 11620	AW575094	EST	7246633
Sequence 11621	AW 576694	EST	7251834
,		·	3 . 3 3 4

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Sequence 11622	AW579819	EST	7254868
Sequence 11623	D44813	EST	1572288
			957166
Sequence 11624	D55269	EST	
Sequence 11625	D80421	EST	1178298
Sequence 11626	F07786	EST	677286
Sequence 11627	F20959	EST	2060135
Sequence 11628	F22113	EST	2061289
Sequence 11629	H01421	EST	864354
			876599
Sequence 11630	H11779	EST	
Sequence 11631	H16460	EST	881280
Sequence 11632	H16546	EST	882771
Sequence 11633	H18560	EST	884800
Sequence 11634	H22094	EST	890789
Sequence 11635	H40387	EST	916439
Sequence 11636	H54240	EST	994387
	H68450	EST	1027190
Sequence 11637			
Sequence 11638	H90819	EST	1081249
Sequence 11639	H94986	EST	1102619
Sequence 11640	L44419	EST	10487 <b>40</b>
Sequence 11641	N20934	EST	1126104
Sequence 11642	N21624	EST	1126794
Sequence 11643	N23545	EST	1137695
	N26941	EST	1141289
Sequence 11644			
Sequence 11645	N28426	EST	1146662
Sequence 11646	N31770	EST	1152169
Sequence 11647	N40973	EST	1164571
Sequence 11648	N44346	EST	1182865
Sequence 11649	N54935	EST	1196255
Sequence 11650	N71715	EST	1228427
Sequence 11651	N80408	EST	1243109
Sequence 11652	R17938	EST	771548
	_	EST	927634
Sequence 11653	R82925		967581
Sequence 11654	R93415	EST	
Sequence 11655	R93745	EST	967911
Sequence 11656	T30702	EST	612800
Sequence 11657	T48489	EST	642689
Sequence 11658	T77803	EST	69 <b>5006</b>
Sequence 11659	T89533	EST	71804 <del>6</del>
Sequence 11660	W01580	EST	1273634
Sequence 11661	W01785	EST	1273784
	W02490	EST	1274488
Sequence 11662			1315016
Sequence 11663	W33011	EST	
Sequence 11664	W79178	EST	1390613
Sequence 11665	V60015	NUCPATENT	N/A
Sequence 11666	V86392	NUCPATENT	N/A
Sequence 11667	X40046	NUCPATENT	N/A
Sequence 11668	X83330	NUCPATENT	949933
Sequence 11669	X87396	NUCPATENT	1304366
•	X97973	NUCPATENT	1360128
Sequence 11670		NUCPATENT	1403131
Sequence 11671	X98409		
Sequence 11672	Z33373	NUCPATENT	534033
Sequence 11673	Z3 <b>3622</b>	NUCPATENT	608977
Sequence 11674	Z33656	NUCPATENT	488829
Sequence 11675	Z39728	NUCPATENT	565523
Sequence 11676	AC02994	PREPATNUC	N/A
Sequence 11677	AC31122	PREPATNUC	N/A
204401106 11077		, , , _ , , , , , , , , , , , , , , , ,	

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Sequence 11678	AC34163	PREPATNUC	N/A
Sequence 11679	AC35230	PREPATNUC	N/A
Sequence 11680	AC36946	PREPATNUC	N/A
Sequence 11681	AC2759	Genbank	
		_	345130
Sequence 11682	A16794	Genbank	512417
Sequence 11683	A17003	Genbank	512419
Sequence 11684	A20255	Genbank	1247401
Sequence 11685	A364 <b>60</b>	Genbank	2293778
Sequence 11686	AB000584	Genbank	1813326
Sequence 11687	AB001535	Genbank	3928755
Sequence 11688	AB002308	Genbank	6634012
Sequence 11689	AB002321	Genbank	
	_		2224586
Sequence 11690	AB002323	Genbank	2224590
Sequence 11691	AB002387	Genbank	2224718
Sequence 11692	AB002390	Genbank	22 <b>8</b> 0487
Sequence 11693	AB002533	Genbank	1944124
Sequence 11694	AB003102	Genbank	1945608
Sequence 11695	AB003151	Genbank	3702678
Sequence 11696	AB003698	Genbank	2102636
Sequence 11697	AB003730	Genbank	2627128
Sequence 11698	AB007166	Genbank	3077761
	AB007882	Genbank	
Sequence 11699	_		2887418
Sequence 11700	AB007887	Genbank	2887428
Sequence 11701	AB007916	Genbank	6683704
Sequence 11702	AB007917	Genbank	3413857
Sequence 11703	AB007922	Genbank	6634036
Sequence 11704	AB007938	Genbank	3413899
Sequence 11705	AB010882	Genbank	2 <del>96</del> 7451
Sequence 11706	AB011004	Genbank	3273315
Sequence 11707	AB011083	Genbank	3043545
Sequence 11708	AB011111	Genbank	3043601
Sequence 11709	AB011139	Genbank	3043657
Sequence 11710	AB011420	Genbank	3834353
•	AB014077		
Sequence 11711		Genbank	5672586
Sequence 11712	AB014554	Genbank	3327121
Sequence 11713	AB015639	Genbank	5821139
Sequence 11714	AB017004	Genbank	4239949
Sequence 11715	AB017103	Genbank	3493334
Sequence 11716	AB017335	Genbank	3582440
Sequence 11717	AB017708	Genbank	3650434
Sequence 11718	AB018260	Genbank	3882154
Sequence 11719	AB018265	Genbank	3882164
Sequence 11720	AB018267	Genbank	3882168
Sequence 11721	AB018304	Genbank	3882242
Sequence 11722	AB018330	Genbank	3882294
Sequence 11723	AB018344		
		Genbank	3882322
Sequence 11724	AB019524	Genbank	4519939
Sequence 11725	AB019568	Genbank	3 <b>88</b> 53 <b>7</b> 1
Sequence 11726	AB020662	Genbank	4240198
Sequence 11727	AB020679	Genbank	4240232
Sequence 11728	AB020684	Genbank	4240242
Sequence 11729	AB020686	Genbank	4240246
Sequence 11730	AB020692	Genbank	4240258
Sequence 11731	ABC20697	Genbank	4240268
Sequence 11732	AB020715	Genbank	
Sequence 11732	AB020858		4240304
Coduence 11733	A0020000	Genbank	4003378

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Sequence	11734	AB020861	Genbank	4003381
Sequence		AB021288	Genbank	4038732
Sequence		AB021467	Genbank	4586518
Sequence		AB021663	Genbank	
_ •				4996450
Sequence		AB022435	Genbank	4996607
Sequence		AB022537	Genbank	6942217
Sequence		AB023050	Genbank	5672605
Sequence:	11741	AB023691	Genbank	6519222
Sequence:	11742	AB023699	Genbank	6519298
Sequence	11743	AB024334	Genbank	6016837
Sequence	11744	AB026894	Genbank	5821423
Sequence :	11745	AB027466	Genbank	6172220
Sequence		AB028859	Genbank	6567165
Sequence		AB029003	Genbank	5689496
Sequence		AB029347	Genbank	5921132
Sequence		AB029551	Genbank	
		AB029337 AB032417		6714542
Sequence :			Genbank	6277265
Sequence		AB032969	Genbank	6329965
Sequence		AB032975	Genbank	6330044
Sequence 1		AB033070	Genbank	6330818
Sequence :		AB033076	Genbank	63 <b>30860</b>
Sequence:		AB033079	Genbank	6382025
Sequence 1	11756	AB033082	Genbank	6330910
Sequence:	11757	AB033117	Genbank	6331435
Sequence :	11758	AB034205	Genbank	6899845
Sequence 1	11759	AB037744	Genbank	7243026
Sequence 1	11760	AB037757	Genbank	7243052
Sequence		AB037763	Genbank	7243064
Sequence		AB037767	Genbank	7243072
Sequence 1		AB037807	Genbank	7243152
Sequence		AB037849	Genbank	7243236
Sequence :		AC000040	Genbank	7923866
Sequence 1		AC000083	Genbank	3810636
Sequence 1		AC000093	Genbank	4678240
Sequence 1		AC000120	Genbank	1809224
Sequence 1		AC000353	Genbank	6970735
Sequence 1		AC000378	Genbank	2270906
Sequence 1		AC000394	Genbank	2133911
Sequence 1		AC001226	Genbank	2133862
Sequence 1		AC002039	Genbank	2342716
Sequence 1		AC002040	Genbank	2347081
Sequence 1		AC002064	Genbank	2076723
Sequence 1		AC002069	Genbank	2076718
Sequence 1	1777	AC002094	Genbank	2155224
Sequence 1	11778	AC002117	Genbank	2281075
Sequence 1	11779	AC002401	Genbank	2599240
Sequence 1	1780	AC002470	Genbank	647 <b>8944</b>
Sequence 1		AC002481	Genbank	2340092
Sequence 1		AC002544	Genbank	3337382
Sequence 1		AC003041	Genbank	3264572
Sequence 1		AC003103	Genbank	2842782
Sequence 1		AC003109	Genbank	2961444
Sequence 1		AC003103	Genbank	2636670
Sequence 1		AC003682	Genbank	3264845
Sequence 1		AC003688	Genbank	3789719
Sequence 1	1789	AC003964	Genbank	298125 <b>3</b>

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<u>Table 3-5</u>

Sequence 11790	ACC03989	Genbank	2772538
Sequence 11791	AC003999	Genbank	2772566
Sequence 11792	AC004006	Genbank	2781390
Sequence 11793	AC004039	Genbank	2811101
Sequence 11794	AC004049	Genbank	3366580
Sequence 11795	AC004050	Genbank	4001528
Sequence 11796	AC004098	Genbank	3097872
Sequence 11797	AC004131	Genbank	3342217
Sequence 11798	AC004142	Genbank	2880078
Sequence 11799	AC004148	Genbank	3482960
Sequence 11800	AC004150	Genbank	4585948
Sequence 11801	AC004156	Genbank	2896799
	AC004130		
Sequence 11802 Sequence 11803	AC004221 AC004228	Genbank Genbank	9665 <b>053</b>
			4263 <b>838</b>
Sequence 11804	AC004258	Genbank	2924759
Sequence 11805	AC004263	Genbank	2935616
Sequence 11806	AC004381	Genbank	2982169
Sequence 11807	AC004408	Genbank	3873185
Sequence 11808	AC004467	Genbank	3132844
Sequence 11809	AC004472	Genbank	2984 <b>582</b>
Sequence 11810	AC004474	Genbank	30 <b>97873</b>
Sequence 11811	AC004509	Genbank	29 <b>96628</b>
Sequence 11812	AC004520	Genbank	30 <b>04572</b>
Sequence 11813	AC004525	Genbank	321 <b>9333</b>
Sequence 11814	AC004527	Genbank	4760 <b>422</b>
Sequence 11815	AC004530	Genbank	31 <b>08050</b>
Sequence 11816	AC004531	Genbank	333 <b>7392</b>
Sequence 11817	AC004552	Genbank	327 <b>3378</b>
Sequence 11818	AC004583	Genbank	329 <b>3210</b>
Sequence 11819	AC004584	Genbank	341 <b>7305</b>
Sequence 11820	AC004643	Genbank	30 <b>9784</b> 1
Sequence 11821	AC004656	Genbank	3253114
Sequence 11822	AC004660	Genbank	310 <b>8018</b>
Sequence 11823	AC004686	Genbank	3 <b>688105</b>
Sequence 11824	AC004742	Genbank	315 <b>2630</b>
Sequence 11825	AC004749	Genbank	315 <b>7918</b>
Sequence 11826	AC004754	Genbank	3165 <b>400</b>
Sequence 11827	AC004771	Genbank	36 <b>68108</b>
Sequence 11828	AC004776	Genbank	3169 <b>299</b>
Sequence 11829	AC004810	Genbank	32 <b>64567</b>
Sequence 11830	AC004812	Genbank	397 <b>0970</b>
Sequence 11831	AC004821	Genbank	475 <b>3291</b>
Sequence 11832	AC004832	Genbank	66241 <b>29</b>
Sequence 11833	AC004839	Genbank	39 <b>8357</b> 1
Sequence 11834	AC004841	Genbank	445 <b>4523</b>
Sequence 11835	AC004846	Genbank	7243 <b>869</b>
Sequence 11836	AC004854	Genbank	482 <b>7328</b>
Sequence 11837	AC004858	Genbank	6624125
Sequence 11838	AC004859	Genbank	50 <b>91656</b>
Sequence 11839	AC004883	Genbank	426 <b>3746</b>
Sequence 11840	AC004890	Genbank	45 <b>08146</b>
Sequence 11841	AC004904	Genbank	4156180
Sequence 11842	AC004912	Genbank	4156178
Sequence 11843	AC004943	Genbank	392 <b>4671</b>
Sequence 11844	AC004950	Genbank	4753 <b>269</b>
Sequence 11845	AC004930 AC004985	Genbank	570 <b>8490</b>
ocquence 11045	A0004303	Genbank	37 <b>00490</b>

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<u>Table 3-5</u>

46 AC0049	90 Gent	oank 3924	668
89 AC0056	27 Genb		
70 AC0056	29 Genb	ank 72438	877
71 AC0056	30 Genb		
		ank 3818	348
73 AC0057	02 Genb	ank 3688 <sup>-</sup>	100
	26 Genb	ank 38106	372
'5 AC0057	86 Genb	ank 37022	287
	89 Genb	ank 37022	281
		ank 38498	320
		ank 39474	427
		ank 4731(	)44
		ank 43144	121
		ank 50916	355
2 AC00628			
1 ACCCTO			
1 AC00/03	sz Genba	ank 55238	132
	47 AC0049 48 AC0050 49 AC0050 50 AC0050 51 AC0050 52 AC0050 53 AC0051 55 AC0051 56 AC0052 57 AC0052 58 AC0053 60 AC0053 61 AC0053 62 AC0053 63 AC0053 64 AC0053 65 AC0054 66 AC0055 67 AC0056 67 AC0056 68 AC0056 67 AC0056 67 AC0056 68 AC0057 67 AC0056 68 AC0057 67 AC0056 68 AC0057 68 AC0057 68 AC0057 68 AC0057 68 AC0058 69 AC0056 69 AC0056 60 AC0060 60 AC0060 60 AC0060 60 AC0060 61 AC0060 62 AC0060 63 AC0060 64 AC0060 65 AC0060 66 AC0060 67 AC0060 67 AC0060 68 AC0060 68 AC0060 69 AC0060 69 AC0060 69 AC0060 60 AC0060 60 AC0060 60 AC0060 61 AC0060 61 AC0060 62 AC0060 63 AC0060 64 AC0060 65 AC0060 66 AC0060 67 AC0060 67 AC0060 68 AC0060 69 AC0060 69 AC0060 60 AC0060	47 AC004999 Genter   48 AC005034 Genter   49 AC005037 Genter   50 AC005041 Genter   51 AC005042 Genter   52 AC005084 Genter   53 AC005089 Genter   54 AC005104 Genter   55 AC005153 Genter   56 AC005210 Genter   57 AC005257 Genter   58 AC005263 Genter   59 AC005324 Genter   50 AC005324 Genter   51 AC005329 Genter   52 AC005332 Genter   53 AC005382 Genter   54 AC005382 Genter   55 AC005488 Genter   56 AC005550 Genter   57 AC005621 Genter   58 AC005621 Genter   58 AC005621 Genter   59 AC005621 Genter   59 AC005621 Genter   50 AC005621 Genter   50 AC005621 Genter   51 AC005621 Genter   52 AC005621 Genter   53 AC005621 Genter   54 AC005621 Genter   55 AC005621 Genter   56 AC005621 Genter   57 AC005629 Genter   58 AC005621 Genter   59 AC005621 Genter   59 AC005630 Genter   50 AC005621 Genter   50 AC005621 Genter   51 AC005630 Genter   52 AC005621 Genter   53 AC005702 Genter   54 AC005629 Genter   55 AC005789 Genter   56 AC005789 Genter   57 AC005837 Genter   58 AC005789 Genter   59 AC005841 Genter   50 AC005841 Genter   50 AC005841 Genter   51 AC005841 Genter   52 AC005841 Genter   53 AC00608 Genter   54 AC00608 Genter   55 AC00608 Genter   56 AC00607 Genter   57 AC006157 Genter   58 AC00607 Genter   58 AC006273 Genter   59 AC006286 Genter   50 AC006286 Ge	47 AC004999 Genbank 9970 48 AC005034 Genbank 9947 49 AC005037 Genbank 4508 50 AC005041 Genbank 4508 51 AC005084 Genbank 3659 52 AC005084 Genbank 5732 53 AC005089 Genbank 4218 54 AC005104 Genbank 6249 55 AC005153 Genbank 3289 56 AC005257 Genbank 3289 57 AC005257 Genbank 3289 58 AC005318 Genbank 3289 59 AC005324 Genbank 3386 50 AC005324 Genbank 3342 51 AC005329 Genbank 3342 52 AC005332 Genbank 3367 53 AC005389 Genbank 3367 54 AC005389 Genbank 3367 56 AC005318 Genbank 3367 57 AC005650 Genbank 3478 58 AC005369 Genbank 3367 59 AC005380 Genbank 3540 50 AC005381 Genbank 3540 51 AC005621 Genbank 3540 52 AC005382 Genbank 3540 53 AC005621 Genbank 3548 54 AC005621 Genbank 3548 55 AC005661 Genbank 3548 56 AC005629 Genbank 7243 57 AC005629 Genbank 3188 57 AC005629 Genbank 3688 58 AC006629 Genbank 47310 58 AC006647 Genbank 47310 58 AC006647 Genbank 47310 58 AC006647 Genbank 47310 58 AC0066480 Genbank 49663 59 AC006566 Genbank 49060 60 AC006581 Genbank 49060

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Sequence 11902	AC007041	Genbank	570 <b>847</b> 1
Sequence 11903	AC007055	Genbank	4885 <b>69</b> 1
Sequence 11904	AC007066	Genbank	4508 <b>098</b>
Sequence 11905	AC007073	Genbank	4416547
Sequence 11906	AC007172	Genbank	4731066
Sequence 11907	AC007191	Genbank	45 <b>58643</b>
Sequence 11908	AC007225	Genbank	6715703
Sequence 11909	AC007253	Genbank	5649375
Sequence 11910	AC007384	Genbank	
Sequence 11911	AC007406	Genbank	57 <b>08473</b> 46 <b>89442</b>
Sequence 11912	AC007528	Genbank	
Sequence 11913	AC007527	Genbank	52 <b>63309</b>
Sequence 11914	AC007541	Genbank	4914348
Sequence 11915	AC007564	Genbank	4982536
Sequence 11916	AC007565	Genbank	4982534
Sequence 11917	AC007736	Genbank	4835751
Sequence 11918	AC007738	Genbank	5732177
Sequence 11919	AC007939	Genbank	5306288
Sequence 11920	AC007999 AC008044		7007656
Sequence 11921	AC008101	Genbank Genbank	6015188
Sequence 11922	AC008101 AC008102		6970743
Sequence 11923	AC008102 AC008122	Genbank Genbank	9797815
Sequence 11924	AC008122 AC008123	Genbank	5931373
Sequence 11925	AC008123 AC008269	Genbank	6006046
Sequence 11926	AC008498	Genbank	7243890
Sequence 11927	AC008498 AC008893	Genbank	685 <b>0299</b>
Sequence 11928	AC009330	Genbank	7259691
Sequence 11929	AC010206	Genbank	655 <b>3960</b>
Sequence 11930	AC010200 AC010582	Genbank	6539154
Sequence 11931	AC010302	Genbank	6721135
Sequence 11932	AC012087	Genbank	992 <b>9686</b> 7114 <b>46</b> 5
Sequence 11933	AC020663	Genbank	6682593
Sequence 11934	AC021049	Genbank	7248920
Sequence 11935	AF001549	Genbank	3355302
Sequence 11936	AF001977	Genbank	2914738
Sequence 11937	AF002668	Genbank	2232173
Sequence 11938	AF003837	Genbank	2228792
Sequence 11939	AF004162	Genbank	304 <b>6385</b>
Sequence 11940	AF004339	Genbank	2245559
Sequence 11941	AF004341	Genbank	2245563
Sequence 11942	AF004342	Genbank	2245565
Sequence 11943	AF005896	Genbank	2209244
Sequence 11944	AF006083	Genbank	2282031
Sequence 11945	AF007544	Genbank	2970122
Sequence 11946	AF007835	Genbank	4102034
Sequence 11947	AF008303	Genbank	2580619
Sequence 11948	AF011889	Genbank	2335186
Sequence 11949	AF012023	Genbank	2305237
Sequence 11950	AF012072	Genbank	9967556
Sequence 11951	AF012126	Genbank	28 <b>32227</b>
Sequence 11952	AF015812	Genbank	259 <b>9359</b>
Sequence 11953	AF016709	Genbank	2731560
Sequence 11954	AF019413	Genbank	2347130
Sequence 11955	AF020057	Genbank	31 <b>50436</b>
Sequence 11956	AF020351	Genbank	265 <b>5052</b>
Sequence 11957	AF020352	Genbank	265 <b>5052</b>
	·		200004

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Sequence 11958	AF021819	Genbank	2460317
Sequence 11959	AF022913	Genbank	2558890
Sequence 11960	AF026292	Genbank	2559009
	AF026816	Genbank	2583078
Sequence 11961			
Sequence 11962	AF028832	Genbank	3287488
Sequence 11963	AF030186	Genbank	3831546
Sequence 11964	AF032387	Genbank	2641556
Sequence 11965	AF033095	Genbank	2645728
Sequence 11966	AF035840	Genbank	3800739
Sequence 11967	AF036715	Genbank	4007385
Sequence 11968	AF038178	Genbank	2795898
Sequence 11969	AF038200	Genbank	2795921
Sequence 11970	AF038554	Genbank	2895558
Sequence 11971	AF038633	Genbank	3252876
	AF038955	Genbank	3329379
Sequence 11972			3493528
Sequence 11973	AF041483	Genbank	
Sequence 11974	AF042166	Genbank	3298596
Sequence 11975	AF042836	Genbank	3776241
Sequence 11976	AF044957	Genbank	4164445
Sequence 11977	AF044958	Genbank	4164447
Sequence 11978	AF047185	Genbank	2909861
Sequence 11979	AF047440	Genbank	3335135
Sequence 11980	AF047472	Genbank	2921872
Sequence 11981	AF049105	Genbank	2984656
Sequence 11982	AF051316	Genbank	6580759
Sequence 11983	AF051894	Genbank	3095110
Sequence 11984	AF052129	Genbank	3360438
Sequence 11985	AF052642	Genbank	2981453
	AF053356	Genbank	3135305
Sequence 11986		Genbank	3005692
Sequence 11987	AF054984		
Sequence 11988	AF054990	Genbank	3005703
Sequence 11989	AF055473	Genbank	nil
Sequence 11990	AF055474	Genbank	nil
Sequence 11991	AF061737	Genbank	4335938
Sequence 11992	AF064019	Genbank	3347856
Sequence 11993	AF064 <b>08</b> 4	Genbank	3135668
Sequence 11994	AF064245	Genbank	3930528
Sequence 11995	AF064861	Genbank	3171154
Sequence 11996	AF067517	Genbank	3850313
Sequence 11997	AF067844	Genbank	4240386
Sequence 11998	AF068754	Genbank	3283408
Sequence 11999	AF069378	Genbank	3982733
Sequence 12000	AF070646	Genbank	3283920
Sequence 12001	AF070649	Genbank	3283923
Sequence 12001	AF070652	Genbank	4454679
			4454685
Sequence 12003	AF070655	Genbank	
Sequence 12004	AF073792	Genbank	3300101
Sequence 12005	AF075053	Genbank	3377594
Sequence 12006	AF075242	Genbank	4377992
Sequence 12007	AF077042	Genbank	4689131
Sequence 12008	AF077043	Genbank	4689133
Sequence 12009	AF077046	Genbank	4689139
Sequence 12010	AF077052	Genbank	4689151
Sequence 12011	AF077202	Genbank	4679017
Sequence 12012	AF078855	Genbank	5531824
Sequence 12013	AF081282	Genbank	4336324
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Sequence 12014	AF083441	Genbank	5813822
Sequence 12015	AF085361	Genbank	5114056
Sequence 12016	AF086168	Genbank	3483513
Sequence 12017	AF086169	Genbank	3483514
Sequence 12018	AF086210	Genbank	3483555
Sequence 12019	AF086236	Genbank	3483581
Sequence 12020	AF086329	Genbank	3483674
•			
Sequence 12021	AF086465	Genbank	3483810
Sequence 12022	AF086557	Genbank	3483902
Sequence 12023	AF091075	Genbank	3859987
Sequence 12024	AF091083	Genbank	3860003
Sequence 12025	AF091084	Genbank	3860005
Sequence 12026	AF092124	Genbank	3659900
Sequence 12027	AF092135	Genbank	5138919
Sequence 12028	AF098162	Genbank	3929582
Sequence 12029	AF098865	Genbank	4204674
Sequence 12030	AF100748	Genbank	5410281
Sequence 12031	AF100928	Genbank	4323586
Sequence 12032	AF103801	Genbank	6048969
Sequence 12033	AF105278	Genbank	4324471
Sequence 12034	AF107893	Genbank	4008128
Sequence 12035	AF109872	Genbank	4028621
Sequence 12036	AF110304	Genbank	7106722
Sequence 12037	AF110460	Genbank	4324698
Sequence 12038	AF110774	Genbank	6523794
Sequence 12039	AF111347	Genbank	7211441
Sequence 12040	AF112215	Genbank	6563217
Sequence 12041	AF112222	Genbank	6563229
Sequence 12042	AF113016	Genbank	6642755
Sequence 12043	AF113680	Genbank	6855605
Sequence 12043	AF113700	Genbank	6855634
Sequence 12045	AF117235	Genbank	6563243
Sequence 12045	AF117829	Genbank	4151947
Sequence 12047	AF118091	Genbank	6650827
		Genbank	
Sequence 12048	AF119663		6563251
Sequence 12049	AF121219	Genbank	4903275
Sequence 12050	AF122004	Genbank	5059061
Sequence 12051	AF128536	Genbank	5305705
Sequence 12052	AF129927	Genbank	4558506
Sequence 12053	AF130249	Genbank	4455134
Sequence 12054	AF131738	Genbank	4406548
Sequence 12055	AF131838	Genbank	4406677
Sequence 12056	AF131856	Genbank	4406702
Sequence 12057	AF132965	Genbank	4680700
Sequence 12058	AF132973	Genbank	4680716
Sequence 12059	AF139065	Genbank	6635330
Sequence 12060	AF144700	Genbank	5107075
Sequence 12061	AF144755	Genbank	5006628
Sequence 12062	AF146191	Genbank	5678818
Sequence 12063	AF147330	Genbank	4761681
Sequence 12064	AF150089	Genbank	5107165
Sequence 12065	AF151020	Genbank	7106761
Sequence 12066	AF151066	Genbank	7106853
Sequence 12067	AF151109	Genbank	6166337
Sequence 12068	AF151807	Genbank	4929566
Sequence 12069	AF151832	Genbank	4929616
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Sequence 12070	AF151841	Genbank	4929634
Sequence 12071	AF151861	Genbank	4929674
Sequence 12072	AF151875	Genbank	4929702
Sequence 12073	AF151879	Genbank	4929710
Sequence 12074	AF151884	Genbank	4929720
Sequence 12075	AF151886	Genbank	4929724
Sequence 12076	AF151889	Genbank	4929730
Sequence 12077	AF151893	Genbank	4929738
Sequence 12077	AF151905	Genbank	4929762
Sequence 12079	AF151906	Genbank	4929764
	AF152462	Genbank	5565976
Sequence 12080	AF155568	Genbank	5031511
Sequence 12081	AF156965	Genbank	5731112
Sequence 12082	* * *	Genbank	
Sequence 12083	AF161389		6841191
Sequence 12084	AF161448	Genbank	6841309
Sequence 12085	AF161466	Genbank	6841455
Sequence 12086	AF161500	Genbank	6841523
Sequence 12087	AF161503	Genbank	6841529
Sequence 12088	AF161512	Genbank	6841547
Sequence 12089	AF161556	Genbank	6841379
Sequence 12090	AF165926	Genbank	5817 <b>858</b>
Sequence 12091	AF168787	Genbank	7239175
Sequence 12092	AF168956	Genbank	5 <b>702387</b>
Sequence 12093	AF1 <b>74496</b>	Genbank	7108914
Sequence 12094	AF176574	Genbank	5 <b>762481</b>
Sequence 12095	AF179274	Genbank	5911 <b>402</b>
Sequence 12096	AF187320	Genbank	6164 <b>847</b>
Sequence 12097	AF187554	Genbank	6 <b>653225</b>
Sequence 12098	AF1 <b>88178</b>	Genbank	6164 <b>864</b>
Sequence 12099	AF188745	Genbank	6425 <b>043</b>
Sequence 12100	AF191018	Genbank	645 <b>7339</b>
Sequence 12101	AF191298	Genbank	7656 <b>642</b>
Sequence 12102	AF196969	Genbank	6289073
Sequence 12103	AF201077	Genbank	645 <b>6748</b>
Sequence 12104	AF203815	Genbank	6979 <b>641</b>
Sequence 12105	AF205588	Genbank	6531 <b>675</b>
Sequence 12106	AF205600	Genbank	6581 <b>094</b>
Sequence 12107	AF207550	Genbank	6470333
Sequence 12108	AF217403	Genbank	6691160
Sequence 12109	AF224669	Genbank	7012905
Sequence 12110	AJ001981	Genbank	254 <b>6978</b>
Sequence 12111	AJ010069	Genbank	3483012
Sequence 12112	AJ010341	Genbank	449 <b>0794</b>
Sequence 12113	AJ011803	Genbank	4753 <b>76</b> 0
Sequence 12114	AJ012499	Genbank	5441359
Sequence 12115	AJ131693	Genbank	4584422
Sequence 12116	AJ224442	Genbank	2911586
Sequence 12117	AJ243669	Genbank	6688156
Sequence 12118	AJ249731	Genbank	5921469
Sequence 12119	AJ250042	Genbank	6013005
Sequence 12120	AJ250915	Genbank	6996445
Sequence 12121	AK000049	Genbank	7019880
Sequence 12122	AK000052	Genbank	70198 <sup>25</sup>
Sequence 12123	AK000068	Genbank	70199
Sequence 12124	AK000083	Genbank	701994
Sequence 12125	AK000171	Genbank	7020084
55446106 12 120	ANOGOTI	GGIDAIIN	, 52000-

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Sequence	12126	AK000221	Genbank	7020163
Sequence	12127	AK000303	Genbank	7020295
Sequence	12128	AK000319	Genbank	7020324
Sequence		AK000395	Genbank	7020455
Sequence		AK000400	Genbank	7020463
Sequence		AK000434	Genbank	7020520
Sequence		AK000462	Genbank	7020566
Sequence		AK000470	Genbank	7020580
Sequence		AK000526	Genbank	7020680
Sequence		AK000560	Genbank	7020000
Sequence		AK000615	Genbank	7020827
Sequence		AK000620	Genbank	7020827
Sequence		AK000624	Genbank	7020839
Sequence		AK000656	Genbank	7020889
Sequence		AK000667	Genbank	7020906
Sequence		AK000669	Genbank	7020900
Sequence		AK000809 AK000818	Genbank	7020909
Sequence		AK000818 AK000931	Genbank	7021126
Sequence		AK000931 AK001067	_	
Sequence		AK001087 AK001088	Genbank Genbank	7022104
		AK001088 AK001163		7022136
Sequence Sequence		AK001193 AK001192	Genbank	7022244
			Genbank	7022292
Sequence		AK001221	Genbank	7022340
Sequence		AK001321	Genbank	7022503
Sequence		AK001338	Genbank	7022533
Sequence		AK001354	Genbank	7022561
Sequence		AK001408	Genbank	7022645
Sequence		AK001441	Genbank	7022699
Sequence		AK001453	Genbank	7022720
Sequence		AK001489	Genbank	7022777
Sequence		AK001510	Genbank	7022808
Sequence		AK001536	Genbank	7022851
Sequence		AK001564 AK001614	Genbank	7022893
Sequence			Genbank	7022977
Sequence		AK001689	Genbank	7023104
Sequence		AK001718	Genbank	7023153
Sequence		AK001764	Genbank	7023235
Sequence		AK001780 AK001902	Genbank	7023265
Sequence			Genbank	7023455
Sequence		AK002031	Genbank	7023667
Sequence		AK002100	Genbank	7023777
Sequence		AL008721	Genbank	3171883
Sequence		AL008723 AL009179	Genbank	5101762
Sequence			Genbank	3217024
Sequence		AL020993	Genbank	3980107
Sequence		AL020997	Genbank	2804155
Sequence		AL021368	Genbank	3080468
Sequence		AL021397	Genbank	3319943
Sequence		AL021707	Genbank	4582132
Sequence		AL021917	Genbank	9367203
Sequence		AL022153	Genbank	3319674
Sequence		AL022238	Genbank	4176442
Sequence		AL022240	Genbank	4826471
Sequence		AL022313	Genbank	4200326
Sequence		AL022326	Genbank	3550039
Sequence	12181	AL022718	Genbank	3763969

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Sequence 12182	AL022725	Genbank	5679748
Sequence 12183	AL023581	Genbank	3445283
Sequence 12184	AL023582	Genbank	3805923
Sequence 12185	AL023803	Genbank	9408726
Sequence 12186	AL023807	Genbank	5679750
Sequence 12187	AL024474	Genbank	3395511
Sequence 12188		Genbank	3287156
Sequence 12189		Genbank	3738133
Sequence 12190		Genbank	3421083
Sequence 12191	AL031295	Genbank	4376011
Sequence 12192		Genbank	4902714
Sequence 12193		Genbank	4835258
Sequence 12194	AL031432	Genbank	4375969
Sequence 12195	AL031665	Genbank	4826504
Sequence 12196	AL031681	Genbank	9408729
Sequence 12197	AL031718	Genbank	5805110
Sequence 12198	AL031774	Genbank	3881948
Sequence 12199	AL033377	Genbank	4826462
Sequence 12200	AL033397	Genbank	4902626
Sequence 12201	AL033517	Genbank	4140315
Sequence 12202	AL033522	Genbank	4191258
Sequence 12203	AL033531	Genbank	6807582
Sequence 12204	AL034548	Genbank	7263904
Sequence 12205	AL035209	Genbank	4160217
Sequence 12206	AL035249	Genbank	4688857
Sequence 12207	AL035398	Genbank	5262352
Sequence 12208	AL035411	Genbank	6136940
Sequence 12209	AL035413	Genbank	6010110
Sequence 12210	AL035454	Genbank	9368364
Sequence 12211	AL035458	Genbank	6624641
Sequence 12212	AL035461	Genbank	5123778
Sequence 12213	AL035541	Genbank	9581764
Sequence 12214	AL035610	Genbank	6249454
Sequence 12215	AL035699	Genbank	4826515
Sequence 12216	AL049280	Genbank	4500037
Sequence 12217	AL049294	Genbank	4500055
Sequence 12218	AL049447	Genbank	4500230
Sequence 12219	AL049461	Genbank	4500250
Sequence 12220	AL049552	Genbank	6010194
Sequence 12221	AL049564	Genbank	4902757
Sequence 12222	AL049646	Genbank	9368462
Sequence 12223	AL049694	Genbank	5650653
Sequence 12224	<b>AL04974</b> 8	Genbank	6572235
Sequence 12225	AL049766	Genbank	5763746
Sequence 12226	AL049779	Genbank	8176894
Sequence 12227	AL049829	Genbank	8217859
Sequence 12228	AL049844	Genbank	5924019
Sequence 12229	AL049849	Genbank	4826529
Sequence 12230	AL049869	Genbank	8217861
Sequence 12231	AL049870	Genbank	7248268
Sequence 12232	AL049911	Genbank	6114591
Sequence 12233	AL049944	Genbank	4884189
Sequence 12234	AL049962	Genbank	4884212
Sequence 12235	AL049964	Genbank	4884214
Sequence 12236	AL049968	Genbank	4884217
Sequence 12237	AL050016	Genbank	4884262

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Sequence 12238	AL050089	Genbank	4884107
Sequence 12239	AL050275	Genbank	4886500
Sequence 12240	AL050277	Genbank	4886454
Sequence 12241	AL050343	Genbank	6137008
Sequence 12242	AL078463	Genbank	5777575
Sequence 12243	AL078621	Genbank	6013067
Sequence 12244	AL079333	Genbank	5738434
Sequence 12245	AL080080	Genbank	5262491
Sequence 12246	AL080102	Genbank	5262526
Sequence 12247	AL080193	Genbank	5262674
Sequence 12248	AL080276	Genbank	5763753
Sequence 12249	AL109672	Genbank	5689836
Sequence 12250	AL109759	Genbank	8176900
Sequence 12251	AL109798	Genbank	64533 <b>86</b>
Sequence 12252	AL109865	Genbank	6911646
Sequence 12253	AL110183	Genbank	5817095
Sequence 12254	AL110185	Genbank	5817098
Sequence 12255	AL110193	Genbank	5817109
Sequence 12256	AL110194	Genbank	5817111
Sequence 12257	AL110202	Genbank	5817121
Sequence 12258	AL110271	Genbank	5817083
Sequence 12259	AL117350	Genbank	7159817
Sequence 12260	AL117450	Genbank	5911900
Sequence 12261	AL117461	Genbank	5911922
Sequence 12262	AL117482	Genbank	5911960
Sequence 12263	AL117648	Genbank	5912240
Sequence 12264	AL117662	Genbank	59122 <b>5</b> 7
Sequence 12265	AL121603	Genbank	6434634
Sequence 12266	AL121652	Genbank	7159615
Sequence 12267	AL121716	Genbank	7159797
Sequence 12268	AL121790	Genbank	8919824
Sequence 12269	AL121823	Genbank	6635879
Sequence 12270	AL121852	Genbank	8176919
Sequence 12271	AL121963	Genbank	7161783
Sequence 12272	AL122036	Genbank	6093222
Sequence 12273	AL122066	Genbank	6102860
Sequence 12274	AL133010	Genbank	6453 <b>416</b>
Sequence 12275	AL133060	Genbank	6453 <b>483</b>
Sequence 12276	AL133069	Genbank	6453 <b>507</b>
Sequence 12277	AL133215	Genbank	7228177
Sequence 12278	AL133216	Genbank	6983 <b>473</b>
Sequence 12279	AL133387	Genbank	7076389
Sequence 12280	AL133453	Genbank	70095 <b>94</b>
Sequence 12281	AL133593	Genbank	6599181
Sequence 12282	AL133662	Genbank	6599 <b>256</b>
Sequence 12283	AL135744	Genbank	6682291
Sequence 12284	AL135959	Genbank	6635874
Sequence 12285	AL136296	Genbank	8217913
Sequence 12286	AL136504	Genbank	7018294
Sequence 12287	AL136831	Genbank	6807 <b>666</b>
Sequence 12288	AL137269	Genbank	68077 <b>03</b>
Sequence 12289	AL137450	Genbank	6808024
Sequence 12290	AL137454	Genbank	6808031
Sequence 12291	AL137481	Genbank	680809 <b>6</b>
Sequence 12292	AL137542	Genbank	6808221
Sequence 12293	AL137673	Genbank	6807841
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<u>Table 3-5</u>

Sequence 12294	AL137714	Genbank	6808128
Sequence 12295	AL137721	Genbank	6808159
Sequence 12296	AL137723	Genbank	6808203
Sequence 12297	AL137818	Genbank	7009598
Sequence 12298	AL157442	Genbank	7018558
Sequence 12299	AL157468	Genbank	7018358
Sequence 12300	AL157493	Genbank	
Sequence 12301	AL158493	Genbank	7018539
Sequence 12302	AL158547	Genbank	7161641
Sequence 12303	AP000008	Genbank	7161742
Sequence 12304	AP000009	Genbank	3132318
Sequence 12305	AP000037	Genbank	4666255
Sequence 12306	AP000038	Genbank	3132347
Sequence 12307	AP000346	Genbank	3132348
Sequence 12308	AP000347	Genbank	5103009
Sequence 12309	AP000473	Genbank	5103010
Sequence 12310	AP000497	Genbank	6942329
Sequence 12311	AP000525	Genbank	5926684
Sequence 12312	AP000547	Genbank	5931503
Sequence 12313	AP000692	Genbank	5931525
Sequence 12314	AP000962	Genbank	6693635
Sequence 12315	AP001063	Genbank	6942330
Sequence 12316	AP001115	Genbank	6693613
Sequence 12317	AP001136	Genbank	6899854
Sequence 12318	AP001137	Genbank Genbank	6970360
Sequence 12319	AP001138	Genbank	6970361
Sequence 12320	AP001172	Genbank	7262569
Sequence 12321	AP001343	Genbank	6983884
Sequence 12322	D00596	Genbank	7209831
Sequence 12323	D13119	Genbank	220135
Sequence 12324	D13318	Genbank	285909
Sequence 12325	D13634	Genbank	286026
Sequence 12326	D14658	Genbank	285992
Sequence 12327	D14665	Genbank	285940 6630617
Sequence 12328	D14694	Genbank	
Sequence 12329	D14696	Genbank	603801 285962
Sequence 12330	D14710	Genbank	559324
Sequence 12331	D16111	Genbank	435637
Sequence 12332	D16947	Genbank	598932
Sequence 12333	D17652	Genbank	409069
Sequence 12334	D23662	Genbank	432362
Sequence 12335	D25274	Genbank	464185
Sequence 12336	D26600	Genbank	565650
Sequence 12337	D29805	Genbank	474986
Sequence 12338	D31885	Genbank	505097
Sequence 12339	D38112	Genbank	644480
Sequence 12340	D38550	Genbank	559708
Sequence 12341	D42052	Genbank	1136740
Sequence 12342	D42063	Genbank	924266
Sequence 12343	D45131	Genbank	1304103
Sequence 12344	D50916	Genbank	1469174
Sequence 12345	D50929	Genbank	1469200
Sequence 12346	D55673	Genbank	870746
Sequence 12347	D63486	Genbank	1469885
Sequence 12348	D78130	Genbank	2443315
Sequence 12349	D79205	Genbank	1754620
			1704020

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Sequence 12350	D79991	Genbank	1136397
Sequence 12351	D80000	Genbank	1136415
Sequence 12352	D83077	Genbank	1304131
Sequence 12353	D83647	Genbank	1209009
Sequence 12354	D83767	Genbank	1913784
Sequence 12355	D84273	Genbank	3894217
Sequence 12356	D84344	Genbank	2340829
Sequence 12357	D86971	Genbank	
Sequence 12358	D86982	Genbank	1504015
Sequence 12359	D87450	Genbank	1504037
			1665788
Sequence 12360	D87666	Genbank	1620016
Sequence 12361	D87684	Genbank	1663703
Sequence 12362	D87742	Genbank	1665824
Sequence 12363	E01932	Genbank	2170180
Sequence 12364	E02339	Genbank	2170574
Sequence 12365	E02628	Genbank	2170856
Sequence 12366	E03413	Genbank	2171629
Sequence 12367	E08204	Genbank	217 <b>632</b> 5
Sequence 12368	E12274	Genbank	3251108
Sequence 12369	E13298	Genbank	3252103
Sequence 12370	E13330	Genbank	3252135
Sequence 12371	E15239	Genbank	5709922
Sequence 12372	J00139	Genbank	182715
Sequence 12373	J01415	Genbank	1944628
Sequence 12374	J02960	Genbank	178203
Sequence 12375	J03537	Genbank	337513
Sequence 12376	J03592	Genbank	339722
Sequence 12377	J03746	Genbank	183655
Sequence 12378	J03779	Genbank	179833
Sequence 12379	J04205	Genbank	178686
Sequence 12380	K00558	Genbank	340020
Sequence 12381	L01057	Genbank	292462
Sequence 12382	L05091	Genbank	388030
Sequence 12383	L05092	Genbank	388031
Sequence 12384	L05628	Genbank	1835658
Sequence 12385	L08125	Genbank	306500
Sequence 12386	L11910	Genbank	292420
Sequence 12387	L12350	Genbank	307505
Sequence 12388	L13806	Genbank	306554
Sequence 12389	L13850	Genbank	292165
Sequence 12390	L14076	Genbank	307437
Sequence 12391	L19185	Genbank	440307
Sequence 12392	L19597	Genbank	306467
Sequence 12393	L20046	Genbank	306741
Sequence 12394	L20941	Genbank	507251
Sequence 12395	L22569	Genbank	348706
Sequence 12396	L25616	Genbank	409465
Sequence 12397	L28809	Genbank	454151
Sequence 12398	L31610	Genbank	1220360
Sequence 12399	L35681	Genbank	532031
Sequence 12400	L38951		
Sequence 12400	L38995	Genbank Genbank	893287
		Genbank	704415
Sequence 12402	L39332	Genbank	1235613
Sequence 12403	L39350	Genbank	1237109
Secuence 12404	L41349	Genbank	762825
Sequence 12405	L43964	Genbank	951202

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Sequence 12406	L47647	Genbank	1000861
Sequence 12407	L76416	Genbank	5566605
Sequence 12408	L76568	Genbank	1905925
Sequence 12409	M11058	Genbank	184243
Sequence 12410	M11167	Genbank	337381
Sequence 12411	M11353	Genbank	184092
Sequence 12412	M13519	Genbank	179461
Sequence 12413	M16342	Genbank	184266
Sequence 12414	M16541	Genbank	1280204
Sequence 12415	M17885	Genbank	190231
Sequence 12416	M17886	Genbank	190233
Sequence 12417	M18930	Genbank	184371
Sequence 12418	M19961	Genbank	180940
Sequence 12419	M20496	Genbank	809235
Sequence 12420	M21142	Genbank	183402
Sequence 12421	M22348	Genbank	190815
Sequence 12422	M22760	Genbank	695359
Sequence 12423	M22919	Genbank	189016
Sequence 12424	M24194	Genbank	187701
Sequence 12425	M24486	Genbank	
•	M24543	Genbank	190785
Sequence 12426		Genbank	341200
Sequence 12427	M25160 M26481	Genbank	189059
Sequence 12428		Genbank	619789
Sequence 12429	M26663		618463
Sequence 12430	M27878	Genbank	1020144
Sequence 12431	M30-148	Genbank Genbank	181154
Sequence 12432 Sequence 12433	M31520 M37583	Genbank	337504
			184059
Sequence 12434	M38188 M58485	Genbank Genbank	189378
Sequence 12435 Sequence 12436			180154
Sequence 12437	M59907	Genbank	189383
	M60457	Genbank	181249
Sequence 12438	M63005 M65209	Genbank Genbank	179981
Sequence 12439 Sequence 12440	M74089	Genbank	184444 182400
		Genbank	
Sequence 12441	M74775		187151
Sequence 12442	M75883	Genbank	432974
Sequence 12443	M77142	Genbank	339700
Sequence 12444	M77836	Genbank	189497
Sequence 12445	M81601	Genbank Genbank	339442
Sequence 12446	M90054 M93036		337579
Sequence 12447		Genbank	182904
Sequence 12448	M96982	Genbank	338262
Sequence 12449	S62140	Genbank	386156
Sequence 12450	S63912	Genbank	399757
Sequence 12451	S82300	Genbank	245388
Sequence 12452	U02031	Genbank	451329
Sequence 12453	U07151	Genbank	460624
Sequence 12454	U09813	Genbank	1008454
Sequence 12455	U09954	Genbank	607792
Sequence 12456	U12128	Genbank	557287
Sequence 12457	U13616	Genbank	608024
Sequence 12458	U14193	Genbank	555899
Sequence 12459	U14972	Genbank	550024
Sequence 12460	U16798	Genbank	806751
Sequence 12461	U17899	Genbank	717053

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<u>Table 3-5</u>

Sequence 12462	U19145	Genbank	914904
Sequence 12463	U21914	Genbank	736410
Sequence 12464	U24223	Genbank	1215670
Sequence 12465	U24231	Genbank	809486
Sequence 12466	U25165	Genbank	887792
Sequence 12467	U25182	Genbank	799380
Sequence 12468	U25372	Genbank	1045393
Sequence 12469	U31657	Genbank	951331
Sequence 12470	U33886	Genbank	1710107
Sequence 12471	U34683	Genbank	1236349
Sequence 12472	U35246	Genbank	1477465
Sequence 12473	U37143	Genbank	1185451
Sequence 12474	U37230	Genbank	1574941
Sequence 12475	U38784	Genbank	1574947
Sequence 12476	U41371	Genbank	1173904
Sequence 12477	U52111	Genbank	8331754
•	U52427	Genbank	1924973
Sequence 12478 Sequence 12479	U54562	Genbank	2351381
		Genbank	
Sequence 12480	U54993	Genbank	4097314
Sequence 12481	U55184	Genbank	4097327
Sequence 12482	U55853	Genbank	2145094 1373420
Sequence 12483	U57847	Genbank	
Sequence 12484	U61167 U65896	•••••	1438934
Sequence 12485		Genbank	1763689
Sequence 12486	U66469	Genbank Genbank	1724072
Sequence 12487	U66871		1519518
Sequence 12488	U68105	Genbank	1562509
Sequence 12489	U68758	Genbank	4097815 1825473
Sequence 12490	U69570	Genbank	
Sequence 12491	U70063	Genbank	1743866
Sequence 12492	U73379	Genbank	2062372
Sequence 12493	U73824	Genbank	1857236
Sequence 12494	U80017	Genbank	1737211
Sequence 12495	U80763	Genbank	2231366
Sequence 12496	U81375	Genbank	1845344
Sequence 12497	U82695	Genbank	6960456
Sequence 12498	U90909	Genbank	1913889
Sequence 12499	U91322	Genbank	2344876
Sequence 12500	U91328	Genbank	2088550
Sequence 12501	U94586	Genbank	1946691
Sequence 12502	U95689	Genbank	2804669
Sequence 12503	X00351	Genbank	28251
Sequence 12504	X02152	Genbank	34312
Sequence 12505	X03083 X06409	Genbank	34309
Sequence 12506		Genbank	30204
Sequence 12507	X15822	Genbank	30146
Sequence 12508	X16064	Genbank	37495
Sequence 12509	X52839	Genbank	36125
Sequence 12510	X54540	Genbank	37457
Sequence 12511	X60221	Genbank	509290
Sequence 12512	X61123	Genbank	29508
Sequence 12513	X64707	Genbank	29382
Sequence 12514	X65923	Genbank	31302
Sequence 12515	X66401	Genbank	34634
Sequence 12516	X66503	Genbank	415848
Sequence 12517	X67247	Genbank	36149

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<u>Table 3-5</u>

Sequence 12518	X68271	Genbank	602341
Sequence 12519	X73114	Genbank	402618
Sequence 12520	X74979	Genbank	400462
Sequence 12521	X75682	Genbank	429094
Sequence 12522	X75861	Genbank	456258
		Genbank	434305
Sequence 12523	X76488		2385366
Sequence 12524	X <b>80</b> 199	Genbank	
Sequence 12525	X81900	Genbank	2274973
Sequence 12526	X84908	Genbank	1502344
Sequence 12527	X98356	Genbank	1648869
Sequence 12528	X <b>995</b> 85	Genbank	1770518
Sequence 12529	Y00815	Genbank	34266
Sequence 12530	Y15286	Genbank	2584788
Sequence 12531	Y17448	Genbank	5002578
Sequence 12532	Z13009	Genbank	31072
Sequence 12533	Z14244	Genbank	30150
Sequence 12534	Z48570	Genbank	695580
Sequence 12535	Z56866	Genbank	1028097
Sequence 12536	Z58044	Genbank	1029275
Sequence 12537	Z68277	Genbank	1130689
Sequence 12538	Z69709	Genbank	1204107
Sequence 12539	Z70233	Genbank	1235542
Sequence 12540	Z73417	Genbank	1322397
Sequence 12541	Z82214	Genbank	5911819
Sequence 12542	Z83313	Genbank	1730462
Sequence 12543	Z83843	Cenbank	2578095
Sequence 12544	Z83848	Genbank	3164072
Sequence 12545	Z83851	Genbank	5441348
Sequence 12546	Z84466	Genbank	3319673
Sequence 12547	Z84812	Genbank	1845164
Sequence 12548	Z85999	Genbank	2326514
Sequence 12549	Z92544	Genbank	1869775
Sequence 12550	Z95126	Genbank	2342581
Sequence 12551	Z97053	Genbank	9650676
Sequence 12552	Z98749	Genbank	4775611
Sequence 12553	Z98950	Genbank	3036783
Sequence 12554	Z99571	Genbank	2791262
Sequence 12555	AA001709	EST	1445523
Sequence 12556	AA002222	EST	1445157
Sequence 12557	AA004427	EST	1448541
Sequence 12558	AA005278	EST	1448740
Sequence 12559	AA007226	EST	1463200
Sequence 12560	AA010519	EST	1471545
Sequence 12561	AA011539	EST	1472565
Sequence 12562	AA018969	EST	1482361
Sequence 12563	AA020756	EST	1485332
Sequence 12564	AA022472	EST	1486571
Sequence 12565	AA024695	EST	1489608
	AA025361	EST	1489375
Sequence 12566 Sequence 12567	AA025400	EST	1491566
•	AA025400 AA025432	EST	1490914
Sequence 12568	AA026215	EST	1492901
Sequence 12569	_		1492323
Sequence 12570	AA026422	EST	1492323
Sequence 12571	AA027229	EST	1494207
Sequence 12572	AA028138	EST	1494207
Sequence 12573	AA029944	EST	1430172

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Secuence 12574	AA035377	EST	15071 <b>01</b>
Sequence 12575	AA036809	EST	1510048
Sequence 12576	AA039705	EST	1515984
Sequence 12577	AA039716	EST	15159 <b>95</b>
Sequence 12578	AA040270	EST	1516675
Sequence 12579	AA040302	EST	1516747
Sequence 12580	AA040863	EST	
Sequence 12581	AA044237	EST	1517159
Sequence 12582	AA044842	EST	1522113
Sequence 12583	AA045223	EST	1523064 1523629
Sequence 12584	AA045780	EST	
Sequence 12585	AA046883	EST	1525739
Sequence 12586	AA046936	EST	1524818
Sequence 12587	AA047046		1524835
Sequence 12588	AA047224	EST	1524944
Sequence 12589	AA047729	EST	1525124
Sequence 12590	AA053607	EST	1527526
Sequence 12591	AA053607 AA054470	EST	1544534
Sequence 12592	AA055504	EST	1545395
Sequence 12593	AA055943	EST	1547843
		EST	1548282
Sequence 12594 Sequence 12595	AA057400 AA059199	EST	1550105
Sequence 12596	AA069597	EST	1552077
	AA074189	EST	1576956
Sequence 12597 Sequence 12598	AA074189 AA074611	EST	1614057
Secmence 12599	AA074611 AA074614	EST	1614677
Sequence 12600	AA075494	EST	1614492
Sequence 12601	AA079690	EST EST	1615364
Sequence 12602	AA081037	EST	1618711 1622972
Sequence 12603	AA082177	EST	1624236
Sequence 12604	AA082640	EST	1624905
Sequence 12605	AA083420	EST	1625481
Sequence 12606	AA084868	EST	1626988
Sequence 12607	AA085763	EST	1629225
Sequence 12608	AA085810	EST	1629342
Sequence 12609	AA088187	EST	1633745
Sequence 12610	AA088758	EST	1634279
Sequence 12611	AA088843	EST	1634355
Sequence 12612	AA091442	EST	1636897
Sequence 12613	AA092589	EST	1637410
Sequence 12614	AA093037	EST	1638242
Sequence 12615	AA093542	EST	1639127
Sequence 12616	AA094831	EST	1640416
Sequence 12617	AA094888	EST	1640465
Sequence 12618	AA095725	EST	1641310
Sequence 12619	AA113354	EST	1665194
Sequence 12620	AA115636	EST	1670771
Sequence 12621	AA120954	EST	1678359
Sequence 12622	AA121266	EST	1678899
Sequence 12623	AA126086	EST	16857 <b>60</b>
Sequence 12624	AA127483	EST	1686755
Sequence 12625	AA127687	EST	1687021
Sequence 12826	AA127718	EST	1687007
Sequence 12627	AA128745	EST	1688653
Sequence 12628	AA132163	EST	1693672
Sequence 12629	AA133361	EST	1690329

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Sequence 12630	AA134275	EST	1691754
Sequence 12631	AA135477	EST	1696553
Sequence 12632	AA135809	EST	1696820
Sequence 12633	AA136084	EST	1697294
Sequence 12000	AA136337	EST	1697545
Sequence 12634		EST	1717155
Sequence 12635	AA147775		1721873
Sequence 12636	AA150361	EST	
Sequence 12637	AA150556	EST	1722069
Sequence 12638	AA151565	EST	1720183
Sequence 12639	AA152225	EST	1721426
Sequence 12640	AA156901	EST	1728581
Sequence 12641	AA166664	EST	1745119
Sequence 12642	AA166955	EST	1745331
Sequence 12643	AA167011	EST	1745386
Sequence 12644	AA167399	EST	1745846
Company 12645	AA169486	EST	1748200
Sequence 12645	AA172003	EST	1751060
Sequence 12646		EST	1753566
Sequence 12647	AA173437	EST	1753883
Sequence 12648	AA173705		1754441
Sequence 12649	AA174161	EST	
Sequence 12650	AA177042	EST	1758180
Sequence 12651	AA178986	EST	1760538
Sequence 12652	AA179059	EST	1760471
Sequence 12653	AA179250	EST	1760619
Sequence 12654	AA179888	EST	1761212
Sequence 12655	AA180349	EST	1761631
Sequence 12656	AA181138	EST	1764605
Sequence 12657	AA182751	EST	1766581
Sequence 12658	AA186328	EST	177 <b>444</b> 6
Sequence 12659	AA187076	EST	1775245
Sequence 12660	AA189093	EST	1776127
Sequence 12661	AA191365	EST	1780231
Carrence 12662	AA191629	EST	1780293
Sequence 12662	AA192253	EST	1782078
Sequence 12663	AA193193	EST	1782646
Sequence 12664		EST	1784830
Sequence 12665	AA195119		1799369
Sequence 12666	AA203650	EST	1799390
Sequence 12667	AA203671	EST	1801826
Sequence 12668	AA206446	EST	1802148
Sequence 12669	AA206761	EST	
Sequence 12670	AA215640	EST	1815385
Sequence 12671	AA216102	EST	1816057
Sequence 12672	AA216292	EST	1816271
Sequence 12673	AA216350	EST	1816315
Sequence 12674	AA223194	EST	1843737
Sequence 12675	AA224460	EST	1845078
Sequence 12676	AA226801	EST	1848430
Sequence 12677	AA232991	EST	1856115
Sequence 12678	AA233294	EST	1856287
Sequence 12679	AA233487	EST	1856517
Sequence 12680	AA233674	EST	185 <b>6667</b>
Sequence 12681	AA236252	EST	1858329
	AA242860	EST	1873678
Sequence 12682	AA243369	EST	1874162
Sequence 12683	AA243646	EST	1874456
Sequence 12684		-	1878762
Sequence 12685	AA248173	EST	10/0/02

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Sequence 12686	AA251702	EST	1886665
Sequence 12687	AA252745	EST	1887715
Sequence 12688	AA255738	EST	1892676
Sequence 12689	AA256348	EST	1891933
Sequence 12690	AA258029	EST	1894479
Sequence 12691	AA258243	EST	1894238
Sequence 12692	AA28C080	EST	1921554
Sequence 12693	AA28C848	EST	
Sequence 12694	AA281553	EST	1923546
Sequence 12695	AA283976		1924231
Sequence 12696	AA284242	EST EST	1928258
Sequence 12697	AA285040		1928542
Sequence 12698	AA285126	EST	1927721
Sequence 12699		EST	1928107
•	AA287500	EST	1933225
Sequence 12700	AA287921	EST	1933744
Sequence 12701	AA287941	EST	1933764
Sequence 12702	AA291316	EST	1939294
Sequence 12703	AA292003	EST	1939980
Sequence 12704	AA292069	EST	1940055
Sequence 12705	AA293612	EST	1941263
Sequence 12706	AA295997	EST	1948334
Sequence 12707	AA298183	EST	1950600
Sequence 12708	AA299920	EST	1952324
Sequence 12709	AA303377	EST	1955711
Sequence 12710	AA307371	EST	1959720
Sequence 12711	AA307611	EST	1959960
Sequence 12712	AA311453	EST	1963779
Sequence 12713	AA312681	EST	1965030
Sequence 12714	AA313693	EST	1966022
Sequence 12715	AA314391	EST	1966740
Sequence 12716	AA316545	EST	1968873
Sequence 12717	AA316727	EST	1969286
Sequence 12718	AA316742	EST	1969141
Sequence 12719	AA316868	EST	1969196
Sequence 12720	<b>AA3</b> 17660	EST	1969986
Sequence 12721	AA318533	EST	1970882
Sequence 12722	<b>AA</b> 318684	EST	1971012
Sequence 12723	AA319233	EST	1971558
Sequence 12724	AA319855	EST	1972181
Sequence 12725	AA327986	EST	1980229
Sequence 12726	AA328203	EST	1980603
Sequence 12727	AA345360	EST	1997595
Sequence 12728	AA346874	EST	1999111
Sequence 12729	AA347532	EST	1999862
Sequence 12730	AA364154	EST	2016503
Sequence 12731	AA364506	EST	2016845
Sequence 12732	AA370825	EST	2023163
Sequence 12733	AA373288	EST	2025893
Sequence 12734	AA380652	EST	2032972
Sequence 12735	AA383018	EST	2035335
Sequence 12736	AA393236	EST	2046205
Sequence 12737	AA393310	EST	2046278
Sequence 12738	AA393525	EST	2046493
Sequence 12739	AA393906	EST	2047017
Sequence 12740	AA398315	EST	2051620
Sequence 12741	AA398413	EST	2051522
400	. 5 .55 5 4 1 5	231	2001022

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<u>Table 3-5</u>

Sequence 12742	AA398612	EST	2051721
Sequence 12743	AA400644	EST	2054515
Sequence 12744	AA401361	EST	2053569
Sequence 12745	AA402539	EST	2056345
Sequence 12746	AA402743	EST	2057061
Sequence 12747	AA402800	EST	2056553
Sequence 12748	AA405045	EST	2063467
Sequence 12749	AA405814	EST	2063797
Sequence 12750	AA406310	EST	2064294
Sequence 12751	AA410936	EST	2070190
Sequence 12752	AA417743	EST	2079544
Sequence 12753	AA418533	EST	2080314
Sequence 12754	AA421297	EST	2100139
Sequence 12755	AA427471	EST	2111333
Sequence 12756	AA427574	EST	2112002
Sequence 12757	AA428516	EST	2112513
Sequence 12757	AA430552	EST	2111127
Sequence 12758	AA430332 AA432282	EST	2114670
Sequence 12759	1A434140	EST	2139054
Sequence 12760	AA434447	EST	2139361
Sequence 12761	AA435671	EST	2140585
Sequence 12762	•	EST	2154255
Sequence 12763	AA442377 AA442786	EST	2155461
Sequence 12764		EST	2155553
Sequence 12765	AA442878 AA447199	EST	2159864
Sequence 12766		EST	2161962
Sequence 12767	AA448292 AA448409	EST	2162079
Sequence 12768	AA446409 AA453183	EST	2166852
Sequence 12769		EST	2167080
Sequence 12770	AA453411	EST	2167416
Sequence 12771	AA453747	EST	2167856
Sequence 12772	AA454187 AA456524	EST	2179100
Sequence 12773	AA458612	EST	2183519
Sequence 12774	AA459528	EST	2184435
Sequence 12775	AA460187	EST	2185003
Sequence 12776	AA460167 AA460266	EST	2185082
Sequence 12777	AA460266 AA463256	EST	2188140
Sequence 12778	AA463899	EST	2188783
Sequence 12779	AA465606	EST	2191773
Sequence 12780	AA467764	EST	2194298
Sequence 12781	AA468230	EST	2194764
Sequence 12782	AA468757	EST	2195291
Sequence 12783	AA470337	EST	2197646
Sequence 12784	AA477608	EST	2206242
Sequence 12785	AA478103	EST	2206737
Sequence 12786	AA478663	EST	2207297
Sequence 12787	AA478669	EST	2207303
Sequence 12788	AA480081	EST	2208232
Sequence 12789	AA483251	EST	2212064
Sequence 12790		EST	2212511
Sequence 12791	AA483698	EST	2216580
Sequence 12792	AA486416 AA491100	EST	2220273
Sequence 12793	AA491505	EST	2218288
Sequence 12794	AA491303 AA493382	EST	2223223
Sequence 12795	AA493382 AA493522	EST	2223363
Sequence 12796			2223959
Sequence 12797	AA494118	EST	2223333

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<u>Table 3-5</u>

Sequence 12798	AA494433	EST	2224220
Sequence 12799	AA496993	EST	2230314
Sequence 12800	AA501429	EST	2236396
Sequence 12801	AA503827	EST	2238794
Sequence 12802	AA505115	EST	2241275
Sequence 12803	AA507605	EST	
			2244044
Sequence 12804	AA507791	EST	2244230
Sequence 12805	AA513367	EST	2251779
Sequence 12806	AA513447	EST	2251859
Sequence 12807	AA514215	EST	2253883
Sequence 12808	AA514331	EST	2253839
Sequence 12809	AA514555	EST	2254155
Sequence 12810	AA514835	EST	2254435
Sequence 12811	AA514972	EST	2254572
Sequence 12812	AA515971	EST	2255571
Sequence 12813	AA521142	EST	2261685
Sequence 12814	AA522471	EST	2263183
Sequence 12815	AA523897	EST	2264825
Sequence 12816	AA524190	EST	2265118
Sequence 12817	AA525960	EST	2268029
Sequence 12818	AA526064	EST	2268133
Sequence 12819	AA526186	EST	2268255
Sequence 12820	AA526891	EST	2268960
Sequence 12821	AA526943	EST	2269012
Sequence 12822	AA527040	EST	2269109
Sequence 12823	AA528775	EST	2270844
Sequence 12824		_	
	AA531540	EST	2274246
Sequence 12825	AA533286	EST	2277382
Sequence 12826	AA533355	EST	2277451
Sequence 12827	AA534137	EST	2278153
Sequence 12828	AA535571	EST	2279824
Sequence 12829	AA541666	EST	2288100
Sequence 12830	AA541813	EST	2288247
Sequence 12831	AA548722	EST	2319004
Sequence 12832	AA551733	EST	2321985
Sequence 12833	AA552169	EST	2322421
Sequence 12834	AA554726	EST	2325265
Sequence 12835	AA555194	EST	2325733
Sequence 12836	AA557191	EST	2327668
Sequence 12837	AA563905	EST	2335544
Sequence 12838	AA564118	EST	2335757
Sequence 12839	AA565885	EST	2337524
Sequence 12840	AA568360	EST	2341414
Sequence 12841	AA568578	EST	2341632
Sequence 12842	AA570171	EST	2344151
Sequence 12843	AA577170	EST	2354644
Sequence 12844	AA577927	EST	2356111
Sequence 12845	AA579876	EST	2355203
Sequence 12846	AA580408	EST	2355735
Sequence 12847	AA582370	EST	2359730
Sequence 12848	AA583169	EST	2367778
Sequence 12849	AA586921	EST	2397735
Sequence 12850	AA587325	EST	2398139
Sequence 12851	AA588161		
Sequence 12852	AA588541	EST	2402336
		EST	2401716
Sequence 12853	AA595339	EST	2410689

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Sequence 12854	AA599217	EST	2432842
Sequence 12855	AA602418	EST	2435914
Sequence 12856	AA604553	EST	2445417
Sequence 12857	AA608603	EST	2457031
Sequence 12858	AA608917	EST	2457345
Sequence 12859	AA608990	EST	2457418
		EST	2459229
Sequence 12860	AA610801		2464625
Sequence 12861	AA613587	EST	2464675
Sequence 12862	AA613637	EST	
Sequence 12863	AA614729	EST	2466925
Sequence 12864	AA626535	EST	2538922
Sequence 12865	AA627229	EST	2540273
Sequence 12866	AA627804	EST	2539899
Sequence 12867	AA627996	EST	2539995
Sequence 12868	AA631218	EST	2553829
Sequence 12869	AA631224	EST	2553835
Sequence 12870	AA631404	EST	2554015
Sequence 12871	AA631684	EST	2554295
Sequence 12872	AA632005	EST	2555419
Sequence 12873	AA632225	EST	2555639
Sequence 12874	AA632328	EST	2555742
Sequence 12875	AA633317	EST	2556731
Sequence 12876	AA633888	EST	2557102
Sequence 12877	AA634120	EST	2557334
Sequence 12878	AA634360	EST	2557574
Sequence 12879	AA635630	EST	2559472
Sequence 12880	AA636029	EST	2559871
Sequence 12881	AA639730	EST	2563509
Sequence 12882	AA640721	EST	2565971
Sequence 12883	AA640779	EST	2566029
Sequence 12884	AA642858	EST	2568076
Sequence 12885	AA643157	EST	2568375
Sequence 12886	AA643611	EST	2568829
Sequence 12887	AA643763	EST	2568981
Sequence 12888	AA649327	EST	2575756
Sequence 12889	AA653201	EST	2589372
Sequence 12890	AA655033	EST	2591187
Sequence 12891	AA657408	EST	2593562
Sequence 12892	AA659388	EST	2595542
Sequence 12893	AA662764	EST	2616755
	AA669846	EST	2631345
Sequence 12894	AA669862	EST	2631361
Sequence 12895		EST	2658643
Sequence 12896	AA678121 AA678246	EST	2658768
Sequence 12897	AA678311	EST	2658833
Sequence 12898			2659218
Sequence 12899	AA678696	EST	2668871
Sequence 12900	AA682980	EST	2669225
Sequence 12901	AA683334	EST	
Sequence 12902	AA687631	EST	2674537
Sequence 12903	AA688365	EST	2675271
Sequence 12904	AA694202	EST	2695140
Sequence 12905	AA699532	EST	2702655
Sequence 12906	AA699712	EST	2702675
Sequence 12907	AA700166	EST	2703129
Sequence 12908	AA701936	EST	2705049
Sequence 12909	AA702410	EST	2705 <b>523</b>

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<u>Table 3-5</u>

Sequence 129	)10 AA702	981	EST	2705994
Sequence 129			EST	2714014
Sequence 129			EST	2715664
Sequence 129			EST	2717106
Sequence 129			EST	2718339
Sequence 129		· · — ·	EST	2727225
Sequence 129			EST	2740971
Sequence 129			EST	2752521
Sequence 129			EST	2752577
Sequence 129			EST	2768114
Sequence 129			EST	2788004
Sequence 129			EST	2788983
Sequence 129			EST	2815972
Sequence 129			EST	2820672
Sequence 129			EST	2821851
Sequence 129			EST	2824063
Sequence 129			EST	2824611
Sequence 129			EST	2834958
Sequence 129			EST	2835158
Sequence 129			EST	2836978
Sequence 129			EST	2837891
Sequence 129			ST	2838107
Sequence 129			EST	2848874
Sequence 129			EST	2875479
Sequence 129			EST	2876454
Sequence 129			EST	2877612
Sequence 129	36 AA809		EST	2878673
Sequence 129	37 AA809	700 E	EST	2879106
Sequence 129		964 E	EST	2879370
Sequence 129	39 AA810			2879479
Sequence 129	40 AA811			2880835
Sequence 129	<b>41</b> AA811	879 E	ST	2881490
Sequence 129	42 AA813	197 E	ST	2 <b>883182</b>
Sequence 129	43 AA814	246 E	ST	2883842
Sequence 129				2 <b>884239</b>
Sequence 129		680 E	ST	28 <b>98992</b>
Sequence 129		003 E	ST	29 <b>00366</b>
Sequence 129				29 <b>01429</b>
Sequence 129				29 <b>02836</b>
Sequence 129				29 <b>05025</b>
Sequence 129				2905047
Sequence 129				289 <b>8667</b>
Sequence 129				2910909
Sequence 129				2912327
Sequence 1295				2929633
Sequence 1295				2930258
Sequence 1295				2 <b>932156</b>
Sequence 1295				2932875
Sequence 1295				2942049
Sequence 1295				2953419
Sequence 1296				2955 <b>709</b>
Sequence 1296				2955 <b>856</b>
Sequence 1296				2957845
Sequence 1296				2968038
Sequence 1296				2969295
Sequence 1296	65 AA8734	+89 E	ST 2	2969611

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<u>Table 3-5</u>

Sequence 12966	AA876033	EST	2984796
Sequence 12967	AA877009	EST	2986086
Sequence 12968	AA877239	EST	2986316
Sequence 12969	AA877769	EST	2986734
Sequence 12970	AA883195	EST	2994671
Sequence 12971	AA884623	EST	2994153
Sequence 12972	AA885297	EST	2994374
Sequence 12973	AA886074	EST	300 <b>1182</b>
Sequence 12974	AA890342	EST	3017221
Sequence 12975	AA894745	EST	3031146
Sequence 12976	AA902974	EST	3038097
Sequence 12977	AA903301	EST	3038424
Sequence 12978	AA910691	EST	3049981
Sequence 12979	AA910881	EST	3050171
Sequence 12980	AA917382	EST	3057272
Sequence 12981	AA931246	EST	3085632
Sequence 12982	AA931567	EST	3085953
Sequence 12983	AA933730	EST	3089 <b>998</b>
Sequence 12984	AA935696	EST	3092853
Sequence 12985	AA937407	EST	3095518
Sequence 12986	AA946596	EST	3109991
Sequence 12987	AA947190	EST	3110585
Sequence 12988	AA948676	EST	3109929
	AA954463		3118158
Sequence 12989		EST EST	
Sequence 12990	AA969495		3144675
Sequence 12991	AA969966	EST	3145479
Sequence 12992	AA972733	EST	3147913
Sequence 12993	AA973170	EST	3148350
Sequence 12994	AA983457	EST	3161982
Sequence 12995	AA987257	EST	3172621
Sequence 12996	AA988988	EST	3174559
Sequence 12997	AA992853	EST	317 <b>9398</b>
Sequence 12998	AA992960	EST	3179505
Sequence 12999	AA995518	EST	3182007
Sequence 13000	Al003314	EST	3203728
Sequence 13001	Al005654	EST	3215164
Sequence 13002	AI016909	EST	3231245
			3232474
Sequence 13003	AI018676	EST	-
Sequence 13004	A1025856	EST	3241469
Sequence 13005	AI032664	EST	3253790
Sequence 13006	A1034460	EST	3255413
Sequence 13007	A1039964	EST	3279158
Sequence 13008	A1040839	EST	3 <b>280033</b>
Sequence 13009	Al041671	EST	3280865
Sequence 13010	AI041813	EST	3 <b>281007</b>
Sequence 13011	AI052135	EST	3308126
Sequence 13012	AI052256	EST	3308247
Sequence 13013	AI052761	EST	3308752
Sequence 13014	AI053744	EST	3321531
Sequence 13015	A1057607	EST	3331473
Sequence 13016	AI061586	EST	6358883
Sequence 13017	AI061642	EST	635 <b>8939</b>
Sequence 13018	AI064691	EST	6358963
Sequence 13019	AI064781	EST	63 <b>59053</b>
Sequence 13020	Al074143	EST	34 <b>00787</b>
Sequence 13021	AI074253	EST	3400897
• -			

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<u>Table 3-5</u>

Sequence 1	3022	AI074350	EST	3400994
Sequence 1		AI075208	EST	3401799
Sequence 1		Al076009	EST	3405187
Sequence 1		A1076793	EST	3405971
Sequence 1		AI079885	EST	3416136
Sequence 1		AIC80676	EST	3416927
Sequence 1		AIC81536	EST	3418328
Sequence 1	3029	AIC86009	EST	3424432
Sequence 1	3030	AIC86963	EST	3425386
Sequence 1	3031	AI087104	EST	3425527
Sequence 1		Al091297	EST	3430356
Sequence 1		AI091340	EST	3430399
Sequence 1		AIC96751	EST	3445859
Sequence 1		AI114651	EST	6359996
Sequence 1		AI123912	EST	3539678
Sequence 1		Al126019	EST	3594533
Sequence 1		AI127959	EST	3596473
Sequence 1		AI133330	EST	6360646
Sequence 1		AI139392	EST	3645364
Sequence 1		Al143894	EST	3665703
Sequence 1:	3042	Al144216	EST	3666025
Sequence 1:	3043	AI146737	EST	3674419
Sequence 13	3044	Al148343	EST	3676025
Sequence 1:	3045	AI159872	EST	3693252
Sequence 1:		AI160110	EST	3693490
Sequence 1		AI168423	EST	3701593
Sequence 1		AI183462	EST	3734100
Sequence 1:		Al184770	EST	3735408
Sequence 1:		Al186840	EST	3737478
Sequence 1		Al187815	EST	3739024
Sequence 1		Al189478	EST	3740687
Sequence 13		Al191342	EST	3742551
Sequence 13		Al198812	EST	3751418
Sequence 1:		AI198882	EST	3751488
Sequence 13		AI202429	EST	3755035
Sequence 13		AI202937	EST	3755543
Sequence 13		Al207682	EST	6361 <b>696</b>
Sequence 13		Al216387	EST	3785428
Sequence 13		Al221087	EST	3803290
Sequence 13		Al240307	EST	38357 <b>0</b> 4
Sequence 13		Al240817	EST	3836214
Sequence 13	3063	Al242075	EST	3837472
Sequence 13	3064	Al244187	EST	3839584
Sequence 13	3065	Al246248	EST	3841645
Sequence 13	3066	Al247411	EST	3842808
Sequence 13		AI248525	EST	3843922
Sequence 13		AI249008	EST	3844405
Sequence 1		AI251793	EST	3848322
Sequence 13		AI253046	EST	3849575
Sequence 13		AI253330	EST	3850451
Sequence 1:		Al253436	EST	3850391
Sequence 13		A1262640	EST	3870843
Sequence 13		A1267506	EST	3886673
Sequence 13		A1267570	EST	3886737
Sequence 13		A1268542	EST	3887709
Sequence 13	3077	Al269205	EST	3888372

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<u>Table 3-5</u>

Sequence 13078	AI275967	EST	3898241
Sequence 13079	Al275998	EST	3898272
Sequence 13080	Al276110	EST	3898384
Sequence 13081	Al278317	EST	3900585
Sequence 13082	Al278766	EST	
			3917000
Sequence 13083	Al278998	EST	3917232
Sequence 13084	Al280044	EST	3918277
Sequence 13085	Al280747	EST	3918980
Sequence 13086	Al281464	EST	3919697
Sequence 13087	Al282246	EST	3920479
Sequence 13088	Al282254	EST	3920487
Sequence 13089	Al285371	EST	3923604
Sequence 13090	AI288305	EST	3930985
Sequence 13091	AI289108	EST	3932372
Sequence 13092	Al290733	EST	3933507
Sequence 13093	Al291613	EST	3934387
Sequence 13094	Al291918	EST	3934692
Sequence 13095	Al302808	EST	3962154
	Al302008	EST	4003999
Sequence 13096			
Sequence 13097	Al335458	EST	4072385
Sequence 13098	Al337195	EST	4074122
Sequence 13099	Al339940	EST	4076867
Sequence 13100	Al341735	EST	4078662
Sequence 13101	Al342244	EST	4079171
Sequence 13102	Al344445	EST	4081651
Sequence 13103	Al345345	EST	4082551
Sequence 13104	Al345677	EST	4082883
Sequence 13105	A1356050	EST	4107671
Sequence 13106	Al361513	EST	4113134
Sequence 13107	Al370743	EST	4149496
Sequence 13108	Al380670	EST	4190523
Sequence 13109	Al381627	EST	4194408
Sequence 13110	Al384017	EST	4196798
Sequence 13111	Al400359	EST	4243446
Sequence 13112	Al420311	EST	4266242
Sequence 13113	Al421401	EST	4267332
Sequence 13114	Al435207	EST	4301566
Sequence 13115	Al446753	EST	4297414
Sequence 13116	A1458730	EST	4311309
Sequence 13117	Al458971	EST	4311550
Sequence 13118	Al459103	EST	4311682
Sequence 13119	AI459338	EST	4311917
Sequence 13120	Al468004	EST	4330094
Sequence 13121	Al469505	EST	4331595
Sequence 13122	Al479751	EST	4372919
Sequence 13123	Al480118	EST	4373286
Sequence 13124	Al492144	EST	4393147
Sequence 13125	Al497917	EST	4389899
	Al499678		4391660
Sequence 13126		EST	
Sequence 13127	AI500313	EST	4392295
Sequence 13128	AI554818	EST	4487181
Sequence 13129	Al559531	EST	4509736
Sequence 13130	A1560688	EST	4511029
Sequence 13131	A1566007	EST	4524459
Sequence 13132	AI567612	EST	4526064
Sequence 13133	AI568146	EST	4526720

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<u>Table 3-5</u>

Sequence 13134	Al568614	EST	4531988
Sequence 13135	Al572 <b>600</b>	EST	4535974
Sequence 13136	Al581048	EST	4565424
Sequence 13137	Al582231	EST	4568128
Sequence 13138	Al588871	EST	4597919
Sequence 13139	Al597782	EST	4606830
Sequence 13140	Al610953	EST	4620120
Sequence 13141	Al611995	EST	4621162
Sequence 13142	Al619754	EST	4628880
Sequence 13143	AI627855	EST	4664655
Sequence 13144	Al631608	EST	4682938
Sequence 13145	Al631996	EST	4683326
Sequence 13146	Al632109	EST	
Sequence 13147	Al632955	EST	4683439
Sequence 13148	Al636794	EST	4684285
Sequence 13149	Al653101	EST	4688124
Sequence 13150	Al653441		4737080
Sequence 13151	Al655644	EST	4737420
Sequence 13152	Al659251	EST	4739623
Sequence 13153	Al659386	EST	4762821
Sequence 13154	Al669708	EST	4762956
Sequence 13155	Al678630	EST	4834482
Sequence 13156		EST	4888812
Sequence 13157	A/681539 A/683912	EST	4891721
Sequence 13158		EST	4894094
Sequence 13159	AI684760	EST	4896054
Sequence 13160	Al685829 Al688152	EST	4897123
Sequence 13161		EST	4899446
Sequence 13162	Al688730	EST	4900024
Sequence 13163	Al690281	EST	4901575
Sequence 13164	Al692476	EST	4969816
Sequence 13165	Al692763	EST	4970103
Sequence 13166	AI694509	EST	4971849
	Ai695805	EST	4983705
Sequence 13167	AI703457	EST	4991357
Sequence 13168	Al741323	EST	5109611
Sequence 13169	A1743098	EST	5111386
Sequence 13170	AI744238	EST	5112526
Sequence 13171	Al744256	EST	5112544
Sequence 13172	AI744984	EST	5113272
Sequence 13173 Sequence 13174	Al749216	EST	5127480
Sequence 13175	Al752250	EST	5130514
	AI753825	EST	5132177
Sequence 13176 Sequence 13177	A1760744	EST	5176411
Sequence 13178	AI762877	EST	5178544
Sequence 13179	Al766072	EST	5232581
Sequence 13180	Al769461	EST	5235970
Sequence 13100	Al792057	EST	5339773
Sequence 13181	AI796688	EST	5362151
Sequence 13182	Al802907	EST	5368368
Sequence 13183	Al803019	EST	5368491
Sequence 13184	A1803566	EST	5369026
Sequence 13185	A1806628	EST	5393194
Sequence 13186	Al811625	EST	5398191
Sequence 13187	Al815404	EST	5430950
Sequence 13188	Al815499	EST	5431045
Sequence 13189	AI81 <b>6161</b>	EST	5431707

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Sequence 13190	AI819909	EST	5438 <b>988</b>
Sequence 13191	Al824193	EST	5444864
•		EST	5447299
Sequence 13192	AI826628		
Sequence 13193	AI859220	EST	5512836
Sequence 13194	Al863349	EST	5527 <b>456</b>
Sequence 13195	Al866561	EST	55306 <b>68</b>
Sequence 13196	Al868330	EST	554134 <b>6</b>
Sequence 13197	Al872857	EST	5546 <b>906</b>
Sequence 13198	Al889718	EST	5594882
Sequence 13199	Al905192	EST	6495 <b>579</b>
Sequence 13200	AI912093	EST	5631948
Sequence 13201	Al921575	EST	5657539
Sequence 13202	Al922278	EST	5658242
	Al925156	EST	5661120
Sequence 13203		EST	5664674
Sequence 13204	Al928710		5671707
Sequence 13205	Al932970	EST	
Sequence 13206	Al939359	EST	5678229
Sequence 13207	Al951970	EST	5744280
Sequence 13208	Al952576	EST	5744886
Sequence 13209	Al954275	EST	5746585
Sequence 13210	Al962421	EST	57 <b>5</b> 51 <b>34</b>
Sequence 13211	Al962746	EST	575545 <b>9</b>
Sequence 13212	Al968255	EST	5765 <b>073</b>
Sequence 13213	Al970812	EST	57 <b>67638</b>
Sequence 13214	Al991063	EST	5837966
Sequence 13215	Al992283	EST	58391 <b>88</b>
Sequence 13216	AL036115	EST	5927646
Sequence 13217	AL036349	EST	5405930
Sequence 13218	AL036413	EST	5405969
Sequence 13219	AL037368	EST	5406772
	AL037505 AL038505	EST	5928459
Sequence 13220		EST	5935315
Sequence 13221	AL040528	EST	5409929
Sequence 13222	AL040985		5421723
Sequence 13223	AL042384	EST	
Sequence 13224	AL044640	EST	5432855
Sequence 13225	AL048351	EST	5936451
Sequence 13226	AL134733	EST	6602920
Sequence 13227	AL157600	EST	7058001
Sequence 13228	AW004978	EST	5853682
Sequence 13229	AW014893	EST	5863650
Sequence 13230	AW020158	EST	5873698
Sequence 13231	AW027717	EST	5886473
Sequence 13232	AW051707	EST	5913989
Sequence 13233	AW069100	EST	6024098
Sequence 13234	AW069166	EST	6024252
Sequence 13235	AW069188	EST	6024274
Sequence 13236	AW070930	EST	6025 <b>928</b>
Sequence 13237	AW083185	EST	6038337
Sequence 13238	AW084353	EST	6039505
Sequence 13239	AW086188	EST	6041262
Sequence 13240	AW088595	EST	6044412
Sequence 13241	AW089333	EST	6046677
Sequence 13242	AW118097	EST	6086681
Sequence 13243	AW136562	EST	6140695
		EST	6196586
Sequence 13244	AW148690	EST	6199832
Sequence 13245	AW151847	E21	0199032

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Sequence	13246	AW166819	EST	6398344
Sequence	13247	AW167129	EST	6398654
Sequence		AW168050	EST	6399575
Sequence		AW178252	EST	6444289
Sequence		AW188275	EST	6462711
Sequence 1		AW188668		
Sequence :			EST	6463028
		AW190301	EST	6464781
Sequence		AW192683	EST	6471382
Sequence		AW194025	EST	6472756
Sequence 1		AW194497	EST	6473323
Sequence 1		AW195584	EST	6474734
Sequence 1		AW196723	EST	6475953
Sequence 1		AW204679	EST	6504151
Sequence 1		AW2C7172	EST	6506668
Sequence 1		AW243325	EST	6577165
Sequence 1		AW2636 <b>82</b>	EST	6640498
Sequence 1		AW2909 <b>69</b>	EST	6697605
Sequence 1		AW292970	EST	6699606
Sequence 1		AW295194	EST	6701830
Sequence 1		AW297219	EST	6703855
Sequence 1	3266	AW301845	EST	6711522
Sequence 1		AW302764	EST	6712444
Sequence 1	3268	AW364986	EST	6869 <b>636</b>
Sequence 1	3269	AW365134	EST	6869784
Sequence 1	3270	AW391931	EST	6896590
Sequence 1		AW402563	EST	6921261
Sequence 1		AW405635	EST	6924692
Sequence 1		AW410936	EST	6936477
Sequence 1		AW411275	EST	6936816
Sequence 1		AW444608	EST	6986370
Sequence 1		AW451259	EST	6992035
Sequence 1		AW474887	EST	7044993
Sequence 1		AW578558	EST	7253607
Sequence 1		F29317	EST	4814943
Sequence 1		F34866	EST	4820492
Sequence 1		F36899	EST	4822525
Sequence 1		H17774	EST	884014
Sequence 1		H54240	EST	994387
Sequence 1		H56010	EST	1004654
Sequence 1		H <b>7</b> 1857	EST	1043673
Sequence 1		H78871	EST	1056960
Sequence 1		H97996	EST	1118881
Sequence 1		N20525	EST	1125480
Sequence 1		N27572	EST	1142053
Sequence 1		N48918		
Sequence 1		N62491	EST EST	1190084 1210320
Sequence 1		N95724	EST	1268009
Sequence 1		R02774		
Sequence 1			EST	752510
		R06525	EST	757145 767467
Sequence 1:		R14391	EST	767467
Sequence 1:		R55374	EST	824669
Sequence 1:		T08334	EST	389362
Sequence 1:		T24823	EST	534448
Sequence 1:		T60891	EST	663928
Sequence 1:		T82957	EST	711245
Sequence 1:	3301	T83814	EST	712102

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Sequence 13302	W01580	EST	1273634
Sequence 13303	W04263	EST	1276162
Sequence 13304	W19343	EST	1295012
Sequence 13305	W23474	EST	1300308
Sequence 13306	W69300	EST	1378561
Sequence 13307	W86630	EST	1400227
Sequence 13308	V21239	NUCPATENT	N/A
Sequence 13309	V34331	NUCPATENT	N/A
Sequence 13310	V43615	NUCPATENT	N/A
Sequence 13310	V84583	NUCPATENT	N/A
Sequence 13311	X03841	NUCPATENT	N/A
Sequence 13312	X30393	NUCPATENT	N/A
Sequence 13313		NUCPATENT	N/A
Sequence 13314	X40045		1488593
Sequence 13315	X84983	NUCPATENT	
Sequence 13316	X97975	NUCPATENT	1340080
Sequence 13317	X98010	NUCPATENT	1360094
Sequence 13318	Z14718	NUCPATENT	6383
Sequence 13319	Z40810	NUCPATENT	566550
Sequence 13320	Z41320	NUCPATENT	567459
Sequence 13321	Z41384	NUCPATENT	567528
Sequence 13322	AC02716	PREPATNUC	N/A
Sequence 13323	AC36924	PREPATNUC	N/A
Sequence 13324	AC36925	PREPATNUC	N/A
Sequence 13325	AC36946	PREPATNUC	N/A
Sequence 13326	AC37777	PREPATNUC	N/A
Sequence 13327	AC38546	PREPATNUC	N/A
Sequence 13328	AC39510	PREPATNUC	N/A
Sequence 13329	A02759	Genbank	345130
Sequence 13330	A06846	Genbank	490055
Sequence 13331	A14133	Genbank	490127
Sequence 13332	A16794	Genbank	512417
Sequence 13333	A16796	Genbank	579569
Sequence 13334	A61385	Genbank	3715797
Sequence 13335	AB000584	Genbank	1813326
Sequence 13336	AB001535	Genbank	3928755
Sequence 13337	AB002303	Genbank	2224550
Sequence 13338	AB002308	Genbank	6634012
Sequence 13339	AB002322	Genbank	6634014
Sequence 13340	AB002323	Genbank	2224590
Sequence 13341	AB002360	Genbank	2224664
Sequence 13342	AB002363	Genbank	2224670
Sequence 13343	AB002364	Genbank	2224672
Sequence 13344	AB002366	Genbank	2224676
Sequence 13345	AB002368	Genbank	2224680
Sequence 13346	AB002387	Genbank	2224718
Sequence 13347	AB002391	Genbank	6683696
Sequence 13348	AB002533	Genbank	1944124
Sequence 13349	AB003102	Genbank	1945608
Sequence 13350	AB003723	Genbank	2911141
Sequence 13351	AB004064	Genbank	6983841
Sequence 13352	AB004574	Genbank	3184394
Sequence 13353	AB004857	Genbank	2911111
Sequence 13354	AB006534	Genbank	2924619
Sequence 13355	AB006626	Genbank	6635126
Sequence 13356	AB007144	Genbank	2911155
Sequence 13357	AB007170	Genbank	3077765

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Sequence 13358	ABC07874	Genbank	2887448
Sequence 13359	ABC07890	Genbank	6634032
Sequence 13360	AB007956	Genbank	3413930
Sequence 13361	AB011083	Genbank	3043545
Sequence 13362	AB011105	Genbank	3043589
Sequence 13363	AB011111	Genbank	3043601
Sequence 13364	AB011120	Gerbank	3043619
Sequence 13365	AB011148	Genbank	3043675
Sequence 13366	AB011149	Genbank	3043677
Sequence 13367	AB011155	Genbank	3043689
Sequence 13368	AB014545	Genbank	3327103
Sequence 13369	AB014547	Genbank	3327107
Sequence 13370	AB014560	Genbank	3327133
Sequence 13371	AB014563	Genbank	3327139
Sequence 13372	AB014570	Genbank	3327153
Sequence 13373	AB014579	Genbank	3327171
Sequence 13374	AB014589	Genbank	3327171
Sequence 13375	AB014600	Genbank	3327213
	AB015639	Genbank	5821139
Sequence 13376	AB015059 AB015752	Genbank	3746106
Sequence 13377	AB013732 AB017103	Genbank	
Sequence 13378		Genbank	3493334
Sequence 13379	AB017335		3582440
Sequence 13380	AB017548	Genbank	3885363
Sequence 13381	AB017654	Genbank	6979002
Sequence 13382	AB017709	Genbank	3650435
Sequence 13383	AB018257	Genbank	3882148
Sequence 13384	AB018260	Genbank	3882154
Sequence 13385	AB018267	Genbank	3882168
Sequence 13386	AB018319	Genbank	3882272
Sequence 13387	AB018330	Genbank	3882294
Sequence 13388	AB018340	Genbank	3882314
Sequence 13389	AB018342	Genbank	3882318
Sequence 13390	AB018344	Genbank	3882322
Sequence 13391	AB018353	Genbank	3882340
Sequence 13392	AB019038	Genbank	6970469
Sequence 13393	AB019408	Genbank	4587126
Sequence 13394	AB019441	Genbank	4512311
Sequence 13395	AB019524	Genbank	4519939
Sequence 13396	AB019527	Genbank	3869126
Sequence 13397	AB019564	Genbank	3885367
Sequence 13398	AB020018	Genbank	6456471
Sequence 13399	AB020236	Genbank	4760601
Sequence 13400	AB020335	Genbank	6518494
Sequence 13401	AB020637	Genbank	4240148
Sequence 13402	AB020656	Genbank	4240186
Sequence 13403	AB020660	Genbank	4240194
Sequence 13404	AB020662	Genbank	4240198
Sequence 13405	AB020676	Genbank	4240226
Sequence 13406	AB020682	Genbank	4240238
Sequence 13407	AB020694	Genbank	4240262
Sequence 13408	AB020705	Genbank	4240284
Sequence 13409	AB020711	Genbank	4240296
Sequence 13410	AB020859	Genbank	4003379
Sequence 13411	AB020868	Genbank	4003 <b>388</b>
Sequence 13412	AB020982	Gerbank	4417209
Sequence 13413	AB021470	Genbank	4586521

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<u>Table 3-5</u>

Sequence 13414	AB021663	Genbank	4996450
Sequence 13415	AB022435	Genbank	4996607
Sequence 13416	AB022537	Genbank	6942217
Sequence 13417	AB022785	Genbank	4210446
Sequence 13418	AB023207	Genbank	4589623
Sequence 13419	AB023208	Genbank	4589625
Sequence 13420	AB023224	Genbank	4589657
Sequence 13421	AB023691	Genbank	6519222
Sequence 13422	AB024334	Genbank	6016837
Sequence 13423	AB025355	Genbank	5103146
Sequence 13424	AB025959	Genbank	6520072
Sequence 13425	AB027466	Genbank	6172220
Sequence 13426	AB028624	Genbank	5103045
Sequence 13427	AB028893	Genbank	6552364
Sequence 13428	AB029020	Genbank	5689530
Sequence 13429	AB029331	Genbank	5295999
Sequence 13430	AB030181	Genbank	6730949
Sequence 13431	AB030905	Genbank	5732186
Sequence 13432	AB032951	Genbank	6329748
Sequence 13433	AB032957	Genbank	6329818
Sequence 13434	AB032962	Genbank	6382013
Sequence 13435	AB032969	Genbank	6329965
Sequence 13436	AB033007	Genbank	6330242
Sequence 13437	AB033018	Genbank	6330350
Sequence 13438	AB033020	Genbank	6330364
Sequence 13439	AB033050	Genbank	6330623
Sequence 13440	AB033067	Genbank	6330797
Sequence 13441	AB033069	Genbank	6330811
Sequence 13442	AB033074	Genbank	6330846
Sequence 13443	AB033076	Genbank	6330860
Sequence 13444	AB033079	Genbank	6382025
Sequence 13445	AB033096	Genbank	6331225
Sequence 13446	AB034205	Genbank	6899845
Sequence 13447	AB037763	Genbank	7243064
Sequence 13448	AB037767	Genbank	7243072
Sequence 13449	AB037807	Genbank	7243152
Sequence 13450	AB037813	Genbank	7243164
Sequence 13451	AC000015	Genbank	4263638
Sequence 13452	AC000025	Genbank	4417318
Sequence 13453	AC000046	Genbank	4731059
Sequence 13454	AC000093	Genbank	4678240
Sequence 13455	AC000120	Genbank	1809224
Sequence 13456	AC000353	Genbank	6970735
Sequence 13457	AC000378	Genbank	2270906
Sequence 13458	AC000394	Genbank	2133911
Sequence 13459	AC001226	Genbank	2133862
Sequence 13460	AC001234	Genbank	4028939
Sequence 13461	AC002038	Genbank	2226439
Sequence 13462	AC002039	Genbank	2342716
Sequence 13463	AC002033 AC002041	Genbank	2576343
Sequence 13464	AC002041	Genbank	2076723
Sequence 13465	AC002081	Genbank	2078453
Sequence 13466	AC002081 AC002094	Genbank	2155224
Sequence 13467	AC002094 AC002096	Genbank	2599243
Sequence 13468	AC002090 AC002347	Genbank	2828783
Sequence 13469	AC002347 AC002350	Genbank	2961443
Sednetice 1940a	MUUU2330	Genbank	2301443

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Secuence 1	3470	AC002365	Genbank	2358015
Sequence 1	3471	AC <b>002366</b>	Gənbank	2739349
Secuence 1		AC002394	Genbank	2815550
Secuence 1		AC <b>002395</b>	Genbank	3540146
Secuence 1		AC002398	Genbank	2529398
Secuence 1				
		AC002401	Genbank	2599240
Sequence 1		AC002487	Genbank	2341013
Sequence 1		AC002522	Genbank	7124014
Sequence 1		AC002542	Genbank	2393733
Sequence 1		AC <b>002544</b>	Genbank	3337382
Sequence 1		AC002558	Genbank	2580474
Sequence 1		AC <b>003004</b>	Genbank	2522284
Sequence 1	3482	AC003007	Genbank	2911728
Sequence 1	3483	AC003026	Genbank	2815549
Sequence 1	3484	AC003029	Genbank	9965520
Sequence 1		AC003037	Genbank	2920805
Sequence 1		AC003041	Genbank	3264572
Sequence 1		AC003046	Genbank	10043285
Sequence 1		AC003070	Genbank	2865212
Sequence 1		AC003103	Genbank	2842782
Sequence 1		AC003103	Genbank	3253131
Sequence 1		AC003663	Genbank	3097871
Sequence 1		AC003682	Genbank	3264845
Sequence 1		AC003688	Genbank	
		AC003963		3789719
Sequence 1			Genbank Genbank	2995604
Sequence 1		AC003989		2772538
		AC004006	Genbank	2781390
Sequence 1		AC004013	Genbank	2781380
Sequence 1		AC004073	Genbank	2822136
Sequence 1		AC004084	Genbank	2822156
Sequence 1:		AC004106	Genbank	3150276
Sequence 1		AC004126	Genbank	3242744
Sequence 1		AC004142	Genbank	2880078
Sequence 1		AC004148	Genbank	3482960
Sequence 1		AC004152	Genbank	2894631
Sequence 1:		AC004156	Genbank	2896799
Sequence 1		AC004158	Genbank	2982170
Sequence 13		AC004221	Genbank	9665053
Sequence 1:		AC004222	Genbank	3108042
Sequence 1		AC004223	Genbank	3253129
Sequence 13		AC004227	Genbank	2911718
Sequence 13		AC004228	Genbank	4263838
Sequence 1:		AC004233	Genbank	2914673
Sequence 13		AC004236	Genbank	2914668
Sequence 13		ACC04258	Genbank	292475 <b>9</b>
Sequence 13		AC004263	Genbank	2935616
Sequence 13	3516	AC004381	Genbank	2982169
Sequence 13		AC004382	Genbank	3252819
Sequence 13		AC004388	Genbank	3046271
Sequence 13	3519	AC004408	Genbank	3873185
Sequence 13	3520	AC004466	Genbank	3617739
Sequence 13		AC004477	Genbank	3688107
Sequence 13	3522	AC004494	Genbank	2996647
Sequence 13		AC004507	Genbank	2996632
Sequence 13		AC004520	Genbank	3004572
Sequence 13		AC004522	Genbank	3006227
	-	<del>-</del>		

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<u>Table 3-5</u>

Sequence 1	3526	AC004527	Genbank	4760422
Sequence 1	3527	AC004531	Genbank	3337392
Sequence 1	3528	AC004539	Genbank	3041851
Sequence 1		AC004582	Genbank	3337307
Sequence 1		AC004584	Genbank	3417305
Sequence 1		AC004585	Genbank	3212882
Sequence 1		AC004587	Genbank	3150014
Sequence 1		AC004600	Genbank	4388746
Sequence 1		AC004601	Genbank	
Sequence 1		AC004611	Genbank	3395446 3080643
Sequence 1		AC004617	Genbank	
Sequence 1		AC004617 AC004622	_	3738114
Sequence 1		AC004622 AC004634	Genbank Genbank	3090700
Sequence 1		AC004634 AC004677		3094993
Sequence 1		AC004677 AC004686	Genbank	3126881
Sequence 1		AC004666 AC004760	Genbank	3688105
			Genbank	3168626
Sequence 1		AC004765	Genbank	4731046
Sequence 1: Sequence 1:		AC004770	Genbank	3212836
		AC004798	Genbank	3184497
Sequence 1:		AC004801	Genbank	4204244
Sequence 1:		AC004812	Genbank	3970970
Sequence 13		AC004813	Genbank	4926912
Sequence 13		AC004832	Genbank	6624129
Sequence 13		AC004835	Genbank	4508154
Sequence 13		AC004842	Genbank	4753286
Sequence 1		AC004846	Genbank	7243869
Sequence 13		AC004854	Genbank	4827328
Sequence 13		AC004876	Genbank	4508148
Sequence 13		AC004890	Genbank	4508146
Sequence 13		AC004904	Genbank	4156180
Sequence 13		AC004908	Genbank	4156179
Sequence 13		AC004938	Genbank	3213059
Sequence 13		AC004941	Genbank	4454517
Sequence 13		AC004950	Genbank	4753269
Sequence 13		AC004955	Genbank	4753267
Sequence 13		AC004969	Genbank	4156159
Sequence 13		AC004982	Genbank	3419846
Sequence 13		AC004983	Genbank	4309885
Sequence 13		AC004985	Genbank	5708490
Sequence 13 Sequence 13		AC004987	Genbank	3213010
•		AC004990	Genbank	3924668
Sequence 13		AC004999	Genbank	3970963
Sequence 13		AC005005	Genbank	4156153
Sequence 13		AC005007	Genbank	4156151
Sequence 13		AC005021	Genbank	3694657
Sequence 13		AC005031	Genbank	3688109
Sequence 13		AC005037	Genbank	4827310
Sequence 13		AC005041	Genbank	4508118
Sequence 13		AC005042	Genbank	4156138
Sequence 13		AC005046	Genbank	6094632
Sequence 13		AC005076	Genbank	4508110
Sequence 13		AC005082	Genbank	5836159
Sequence 13		AC005084	Genbank	3659503
Sequence 13		AC005088	Genbank	4753221
Sequence 13		AC005095	Genbank	4753220
Sequence 13	1581	AC005104	Genbank	4218027

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# **Table 3-5**

Sequence 13582	AC005166	Genbank	3242769
Sequence 13583	AC005187	Genbank	3522927
Sequence 13584	AC005189	Genbank	3264580
Sequence 13585	AC005193	Genbank	3309101
Sequence 13586	AC005203	Genbank	3273381
Sequence 13587	AC005212	Genbank	3287444
Sequence 13588	AC005251	Genbank	3287718
Sequence 13589	AC005253	Genbank	3288886
Sequence 13590	AC005261	Genbank	3289984
Sequence 13591	AC005262	Genbank	3289981
Sequence 13592	AC005300	Genbank	6492490
Sequence 13593	AC005318	Genbank	
Sequence 13594	AC005332	Genbank	3885345
Sequence 13595	AC005360	Genbank	3659494
Sequence 13596	AC005363		3367514
Sequence 13597	AC005366	Genbank	3367511
•		Genbank	3367508
Sequence 13598	AC005387	Genbank	3395417
Sequence 13599	AC005391	Genbank	3399673
Sequence 13600	AC005406	Genbank	4580403
Sequence 13601	AC005409	Genbank	4249432
Sequence 13602	AC005410	Genbank	3450753
Sequence 13603	AC005484	Genbank	5091654
Sequence 13604	AC005488	Genbank	5836194
Sequence 13605	AC005509	Genbank	4176355
Sequence 13606	AC005531	Genbank	4153875
Sequence 13607	AC005550	Genbank	3478668
Sequence 13608	AC005562	Genbank	4153858
Sequence 13609	AC005613	Genbank	354 <b>0152</b>
Sequence 13610	AC005627	Genbank	7243867
Sequence 13611	AC005629	Genbank	7243 <b>877</b>
Sequence 13612	AC005630	Genbank	41598 <b>82</b>
Sequence 13613	AC005661	Genbank	3818348
Sequence 13614	AC005668	Genbank	3641771
Sequence 13615	AC005669	Genbank	37023 <b>56</b>
Sequence 13616	AC005726	Genbank	3810672
Sequence 13617	AC005730	Genbank	3779042
Sequence 13618	AC005740	Genbank	3687210
Sequence 13619	AC005768	Genbank	6598827
Sequence 13620	AC005785	Genbank	3702290
Sequence 13621	AC005789	Genbank	3702281
Sequence 13622	AC005828	Genbank	37 <b>89713</b>
Sequence 13623	AC005829	Genbank	3873 <b>300</b>
Sequence 13624	AC005837	Genbank	3849 <b>820</b>
Sequence 13625	AC005838	Genbank	3947427
Sequence 13626	AC005839	Genbank	4079626
Sequence 13627	AC005841	Genbank	4731044
Sequence 13628	AC005876	Genbank	624 <b>9672</b>
Sequence 13629	AC005880	Genbank	624 <b>9667</b>
Sequence 13630	AC005899	Genbank	3935221
Sequence 13631	AC005912	Genbank	4165005
Sequence 13632	AC005923	Genbank	4309927
Sequence 13633	AC005924	Genbank	4309926
Sequence 13634	AC005969	Genbank	4309946
Sequence 13635	AC005998	Genbank	5649387
Sequence 13636	AC006009	Genbank	4753278
Sequence 13637	AC006011	Genbank	4508143

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Sequence 13638	AC006017	Genbank	4508141
Sequence 13639	AC006023	Genbank	4753260
Sequence 13640	AC006026	Genbank	6042101
Sequence 13641	AC006027	Genbank	4156149
Sequence 13642	AC006050	Genbank	4079627
Sequence 13643	AC006057	Genbank	473104 <b>8</b>
Sequence 13644	AC006064	Genbank	4572650
Sequence 13645	AC006084	Genbank	3953485
Sequence 13646	AC006088	Genbank	4204701
Sequence 13647	AC006129	Genbank	3970933
Sequence 13648	AC006137	Genbank	9887710
Sequence 13649	AC006141	Genbank	4580411
Sequence 13650	AC006153	Genbank	4309917
Sequence 13651	AC006160	Genbank	5701616
Sequence 13652	AC006205	Genbank	5815508
Sequence 13653	AC006206	Genbank	4309925
Sequence 13654	AC006207	Genbank	4309924
Sequence 13655	AC006238	Genbank	4204704
Sequence 13656	AC006251	Genbank	4309945
Sequence 13657	AC006254	Genbank	4726099
Sequence 13658	AC006255	Genbank	5306223
Sequence 13659	AC006257	Genbank	4092478
Sequence 13660	AC006273	Genbank	4096060
Sequence 13661	AC006347	Genbank	4309919
Sequence 13662	AC006349	Genbank	7243873
Sequence 13663	AC006409	Genbank	4454486
Sequence 13664	AC006430	Genbank	6453929
Sequence 13665	AC006449	Genbank	61026 <b>66</b>
Sequence 13666	AC006450	Genbank	6997 <b>336</b>
Sequence 13667	AC006475	Genbank	4753283
Sequence 13668	AC006480	Genbank	6289252
Sequence 13669 Sequence 13670	AC006487	Genbank	4314418
Sequence 13671	AC006504 AC006509	Genbank	4220442
Sequence 13672		Genbank	5815507
Sequence 13673	AC006512 AC006518	Genbank	4926863
Sequence 13674	AC006516 AC006536	Genbank Genbank	4713939
Sequence 13675	AC006539	Genbank	4235135
Sequence 13676	AC006561	Genbank	4235141
Sequence 13677	AC006581	Genbank	4558534 4914350
Sequence 13678	AC006597	Genbank	4512725
Sequence 13679	AC006930	Genbank	4309680
Sequence 13680	AC006973	Genbank	4699968
Sequence 13681	AC006987	Genbank	5123989
Sequence 13682	AC007014	Genbank	4371263
Sequence 13683	AC007029	Genbank	7243872
Sequence 13684	AC007038	Genbank	5708475
Sequence 13685	AC007041	Genbank	5708471
Sequence 13686	AC007050	Genbank	6456174
Sequence 13687	AC007055	Genbank	4885691
Sequence 13688	AC007057	Genbank	4680440
Sequence 13689	AC007114	Genbank	4581193
Sequence 13690	AC007130	Genbank	5001538
Sequence 13691	AC007172	Genbank	4731066
Sequence 13692	AC007204	Genbank	4559317
Sequence 13693	AC007207	Genbank	6466489

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### **Table 3-5**

Sequence 13694	AC007216	Genbank	6806842
Sequence 13695	AC007225	Genbank	6715703
Sequence 13696	AC007229	Genbank	4567173
Sequence 13697	AC007239	Genbank	5 <b>523828</b>
Sequence 13698	AC007252	Genbank	5091646
Sequence 13699	AC007314	Genbank	4662678
Sequence 13700	AC007324	Genbank	7923342
Sequence 13701	AC007327	Genbank	4587630
Sequence 13701	AC007327	Genbank	6358845
Sequence 13702	AC007393	Genbank	5306300
Sequence 13703	AC007333	Genbank	5069495
	AC007440 AC007528	Genbank	5263309
Sequence 13705	AC007526 AC007536	Genbank	5917929
Sequence 13706		Genbank	4982536
Sequence 13707	AC007541	Genbank	5668756
Sequence 13708	AC007546		4809347
Sequence 13709	AC007551	Genbank	
Sequence 13710	AC007566	Genbank	4835815
Sequence 13711	AC007766	Genbank	5030437
Sequence 13712	AC007848	Genbank	5923645
Sequence 13713	AC007860	Genbank	5649179
Sequence 13714	AC007868	Genbank	6041758
Sequence 13715	AC007934	Genbank	6721208
Sequence 13716	AC008009	Genbank	566 <b>8757</b>
Sequence 13717	AC008018	Genbank	795 <b>897</b> 3
Sequence 13718	AC008102	Genbank	9 <b>797815</b>
Sequence 13719	AC008115	Genbank	5801 <b>65</b> 4
Sequence 13720	AC008123	Genbank	6006046
Sequence 13721	AC008151	Genbank	5 <b>629925</b>
Sequence 13722	AC008498	Genbank	6850299
Sequence 13723	AC008545	Genbank	6721145
Sequence 13724	AC008583	Genbank	715 <b>889</b> 4
Sequence 13725	AC008893	Genbank	7259691
Sequence 13726	AC008981	Genbank	6721142
Sequence 13727	AC009298	Genbank	7243882
Sequence 13728	AC009363	Genbank	6648145
Sequence 13729	AC009509	Genbank	6492473
Sequence 13730	AC009946	Genbank	6604542
Sequence 13731	AC010077	Genbank	5870275
Sequence 13732	AC010168	Genbank	6855156
Sequence 13733	AC010197	Genbank	6143841
Sequence 13734	AC010436	Genbank	6938861
Sequence 13735	AC010525	Genbank	7248932
Sequence 13736	AC010582	Genbank	6721135
Sequence 13737	AC010722	Genbank	6587939
Sequence 13738	AC011331	Genbank	9929686
Sequence 13739	AC011599	Genbank	7113871
Sequence 13740	AC012085	Genbank	6634755
Sequence 13741	AC012087	Genbank	7114465
Sequence 13742	AC016831	Genbank	6539289
Sequence 13743	AC016941	Genbank	8567654
Sequence 13744	AC020663	Genbank	6682593
	AC020663 AC021049	Genbank	7248920
Sequence 13745		Genbank	2580 <b>549</b>
Sequence 13746	AF000982	Genbank	3355302
Sequence 13747	AF001549	Genbank	2339976
Sequence 13748	AF001552		2232173
Sequence 13749	AF002668	Genbank	22321/3

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Sequence 13750	AF004162	Genbank	3046385
Sequence 13751	AF004337	Genbank	2245557
Sequence 13752	AF004338	Genbank	2245558
Sequence 13753	AF004339	Genbank	2245559
Sequence 13754	AF004342	Genbank	2245565
Sequence 13755	AF006513	Genbank	2645428
Sequence 13756	AF006621	Genbank	7629276
Sequence 13757	AF007134	Genbank	2852608
Sequence 13758	AF007146	Genbank	2852624
Sequence 13759	AF007189	Genbank	2459927
Sequence 13760	AF007544	Genbank	2970122
Sequence 13761	AF007872	Genbank	2358280
Sequence 13762	AF007875	Genbank	
Sequence 13763	AF008303	Genbank	2258417
Sequence 13764	AF009242		2580619
Sequence 13765	AF010313	Genbank	2338289
Sequence 13766	AF011889	Genbank Genbank	6468761
Sequence 13767	AF013488		2335186
Sequence 13768	AF013758	Genbank	2343184
Sequence 13769		Genbank	3046899
Sequence 13770	AF013970 AF014398	Genbank	2801421
Sequence 13771		Genbank	8148061
	AF015812	Genbank	2599359
Sequence 13772	AF017790	Genbank	2501872
Sequence 13773 Sequence 13774	AF020352	Genbank	2655054
	AF020782	Genbank	2522480
Sequence 13775	AF021819	Genbank	2460317
Sequence 13776 Sequence 13777	AF022211	Genbank	2465723
Sequence 13778	AF022728	Genbank	2935182
Sequence 13779	AF026802	Genbank	5712085
Sequence 13779	AF026852	Genbank	3599965
Sequence 13781	AF027205 AF028832	Genbank	2598967
Sequence 13782	AF029750	Genbank Genbank	3287488
Sequence 13783	AF029786	Genbank	2587057
Sequence 13784	AF031379		3403166
Sequence 13785	AF035286	Genbank Genbank	4894208
Sequence 13786	AF035296	Genbank	2661038
Sequence 13787	AF035304	Genbank	2661052
Sequence 13788	AF035840	Genbank	2661064
Sequence 13789	AF036715	Genbank	3800739
Sequence 13790	AF036718	Genbank	4007385
Sequence 13791	AF037447	Genbank	2708629
Sequence 13792	AF038200	Genbank	6466790
Sequence 13793	AF038406	Genbank	2795921
Sequence 13794	AF039019	Genbank	3523122 2828109
Sequence 13795	AF039575	Genbank	
Sequence 13796	AF039594	Genbank	2773157
Sequence 13797	AF040628	Genbank	3366667 3869269
Sequence 13798	AF042836	Genbank	
Sequence 13799	AF043105	Genbank	3776241 3169541
Sequence 13800	AF044956	Genbank	
Sequence 13801	AF044957	Genbank	5326827
Sequence 13802	AF044959	Genbank	4164445
Sequence 13803	AF045447	Genbank	3348136
Sequence 13804	AF047184	Genbank Genbank	2865655
Sequence 13805	AF047185	Genbank	2909859
= = ===================================	07/100	Gennank	2909861

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<u>Table 3-5</u>

Sequence 13806	AF047438	Genbank	3335131
Sequence 13807	AF047440	Genbank	3335135
Sequence 13808	AF047472	Genbank	2921872
Sequence 13809	AF050638	Genbank	5326819
Sequence 13810	AF050639	Genbank	4164453
Sequence 13811	AF052092	Genbank	3360338
Sequence 13812	AF052129	Genbank	3360438
Sequence 13813	AF052164	Genbank	3360475
Sequence 13814	AF052182	Genbank	3360494
Sequence 13815	AF052573	Genbank	3510694
Sequence 13816	AF052578	Genbank	2967847
Sequence 13817	AF053356	Genbank	3135305
Sequence 13818	AF054183	Genbank	4092053
Sequence 13819	AF054990	Genbank	3005703
	AF055066		
Sequence 13820 Sequence 13821	AF055474	Genbank Genbank	3273727
			nil
Sequence 13822	AF055584	Genbank	3649607
Sequence 13823	AF057356	Genbank	3063652
Sequence 13824	AF057735	Genbank	4165042
Sequence 13825	AF058293	Genbank	3047377
Sequence 13826	AF058988	Genbank	3300089
Sequence 13827	AF059527	Genbank	4454661
Sequence 13828	AF060798	Genbank	3372665
Sequence 13829	AF061737	Genbank	4335938
Sequence 13830	AF062077	Genbank	3641524
Sequence 13831	AF064019	Genbank	3347856
Sequence 13832	AF064861	Genbank	3171154
Sequence 13833	AF067139	Genbank	3337440
Sequence 13834	AF068302	Genbank	4584876
Sequence 13835	AF068754	Genbank	3283408
Sequence 13836	AF068755	Genbank	4321979
Sequence 13837	AF069378	Genbank	3 <b>982733</b>
Sequence 13838	AF069532	Genbank	3892190
Sequence 13839	AF070523	Genbank	3764088
Sequence 13840	AF070539	Genbank	3387896
Sequence 13841	AF070556	Genbank	3387921
Sequence 13842	AF070630	Genbank	3283898
Sequence 13843	AF070635	Genbank	3283905
Sequence 13844	AF070638	Genbank	3283909
Sequence 13845	AF070646	Genbank	32 <b>83920</b>
Sequence 13846	AF070649	Genbank	32 <b>83923</b>
Sequence 13847	AF070655	Genbank	4454685
Sequence 13848	AF070659	Genbank	445 <b>4693</b>
Sequence 13849	AF070669	Genbank	4454713
Sequence 13850	AF070673	Genbank	3978241
Sequence 13851	AF072097	Genbank	5725511
Sequence 13852	AF073298	Genbank	3641537
Sequence 13853	AF075061	Genbank	3377602
Sequence 13854	AF077030	Genbank	4689107
Sequence 13855	AF077036	Genbank	4689119
Sequence 13856	AF077043	Genbank	4689133
Sequence 13857	AF077044	Genbank	4689135
Sequence 13858	AF077052	Genbank	4689151
Sequence 13859	AF077200	Genbank	4679013
Sequence 13860	AF077202	Genbank	467 <b>9017</b>
Sequence 13861	AF078847	Genbank	5531 <b>808</b>

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Sequence 13862	AF078850	Genbank	5531814
Sequence 13863	AF082283	Genbank	4092066
Sequence 13864	AF082513	Genbank	3649656
Sequence 13865	AF083441	Genbank	5813822
Sequence 13866	AF085844	Genbank	3483158
Sequence 13867	AF086210	Genbank	3483555
Sequence 13868	AF086329	Genbank	3483674
Sequence 13869	AF086337	Genbank	3483682
Sequence 13870	AF086378	Genbank	3483723
Sequence 13871	AF086490	Genbank	3483835
Sequence 13872	AF086557	Genbank	3483902
Sequence 13873	AF086564	Genbank	3483909
Sequence 13874	AF086628	Genbank	4240457
Sequence 13875	AF087020	Genbank	3851144
Sequence 13876	AF087160	Genbank	3869380
Sequence 13877	AF087959	Genbank	3523165
Sequence 13878	AF088060	Genbank	3523266
Sequence 13879	AF088071	Genbank	3523277
Sequence 13880	AF090095	Genbank	4063630
Sequence 13881	AF090947	Genbank	6690255
Sequence 13882	AF091075	Genbank	3859987
Sequence 13883	AF092094	Genbank	4426606
Sequence 13884	AF092124	Genbank	3659900
Sequence 13885	AF092128	Genbank	5138905
Sequence 13886	AF092130	Genbank	5138909
Sequence 13887	AF092135	Genbank	51 <b>38919</b>
Sequence 13888	AF097026	Genbank	6409120
Sequence 13889	AF097514	Genbank	480 <b>8600</b>
Sequence 13890	AF098162	Genbank	3929582
Sequence 13891	AF100615	Genbank	4808624
Sequence 13892	AF100616	Genbank	4808626
Sequence 13893	AF100748	Genbank	541 <b>0281</b>
Sequence 13894	AF100757	Genbank	5410299
Sequence 13895 Sequence 13896	AF100760	Genbank	5410305
Sequence 13897	AF102803 AF104233	Genbank	4092759
Sequence 13898	AF104913	Genbank	5880481
Sequence 13899	AF106622	Genbank	3941723
Sequence 13900	AF106681	Genbank	4378528
Sequence 13901	AF109134	Genbank Genbank	5410327
Sequence 13902	AF109718	Genbank	4139227
Sequence 13903	AF110460	Genbank	3978547
Sequence 13904	AF111847	Genbank	4324698 7211441
Sequence 13905	AF112213	Genbank	6563213
Sequence 13906	AF112222	Genbank	6563229
Sequence 13907	AF112227	Genbank	4545218
Sequence 13908	AF113011	Genbank	6642745
Sequence 13909	AF113016	Genbank	6642755
Sequence 13910	AF113141	Genbank	4512031
Sequence 13911	AF113535	Genbank	6523824
Sequence 13912	AF113540	Genbank	6523834
Sequence 13913	AF113615	Genbank	5106955
Sequence 13914	AF113680	Genbank	6855605
Sequence 13915	AF113690	Genbank	6855619
Sequence 13916	AF113700	Genbank	685 <b>563</b> 4
Sequence 13917	AF113701	Genbank	6855635
			200000

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<u>Table 3-5</u>

Sequence 13918	AF115402	Genbank	4559272
Sequence 13919	AF116910	Genbank	476 <b>8837</b>
Sequence 13920	AF117235	Genbank	6563243
Sequence 13921	AF117888	Genbank	57 <b>32617</b>
Sequence 13922	AF118084	Genbank	6650813
Sequence 13923	AF118124	Genbank	4235636
Sequence 13924	AF118649	Genbank	684 <b>0946</b>
Sequence 13925	AF119297	Genbank	4633508
Sequence 13926	AF123052	Genbank	5106556
Sequence 13927	AF125530	Genbank	6563 <b>259</b>
Sequence 13928	AF126023	Genbank	6563275
Sequence 13929	AF126782	Genbank	6318547
Sequence 13930	AF127577	Genbank	7019596
Sequence 13931	AF128527	Genbank	4928043
Sequence 13932	AF128536	Genbank	5305705
Sequence 13933	AF129756	Genbank	4337095
Sequence 13934	AF131738	Genbank	4406548
Sequence 13935	AF131748	Genbank	4406563
Sequence 13936	AF131758	Genbank	4406576
Sequence 13937	AF131771	Genbank	44065 <b>95</b>
Sequence 13938	AF131777	Genbank	4406602
Sequence 13939	AF131781	Genbank	4406608
Sequence 13940	AF131814	Genbank	4406 <b>648</b>
Sequence 13941	AF131838	Genbank	4406677
Sequence 13942	AF131856	Genbank	44067 <b>02</b>
Sequence 13943	AF132939	Genbank	468 <b>0648</b>
Sequence 13944	AF132942	Genbank	468 <b>0654</b>
Sequence 13945	AF132973	Genbank	468071 <b>6</b>
Sequence 13946	AF132984	Genbank	495 <b>9567</b>
Sequence 13947	AF134726	Genbank	452 <b>9886</b>
Sequence 13948	AF135162	Genbank	7259481
Sequence 13949	AF135488	Genbank	566 <b>8619</b>
Sequence 13950	AF146651	Genbank	502 <b>0073</b>
Sequence 13951	AF147330	Genbank	4761 <b>681</b>
Sequence 13952	AF147378	Genbank	4761 <b>729</b>
Sequence 13953	AF147412	Genbank	476 <b>1763</b>
Sequence 13954	AF150089	Genbank	510 <b>7165</b>
Sequence 13955	AF151052	Genbank	7106 <b>825</b>
Sequence 13956	AF151103	Genbank	575 <b>8136</b>
Sequence 13957	AF151109	Genbank	616 <b>633</b> 7
Sequence 13958	AF151801	Genbank	49 <b>29554</b>
Sequence 13959	AF151807	Genbank	4929 <b>566</b>
Sequence 13960	AF151832	Genbank	4929 <b>616</b>
Sequence 13961	AF151835	Genbank	492 <b>9622</b>
Sequence 13962	AF151878	Genbank	492 <b>9708</b>
Sequence 13963	AF151903	Genbank	4929758
Sequence 13964	AF151905	Genbank	4929 <b>762</b>
Sequence 13965	AF151906	Genbank	492 <b>9764</b>
Sequence 13966	AF152363	Genbank	56 <b>69134</b>
Sequence 13967	AF153273	Genbank	5932030
Sequence 13968	AF153274	Genbank	5932031
Sequence 13969	AF153608	Genbank	5231140
Sequence 13970	AF156965	Genbank	5731112
Sequence 13971	AF158180	Genbank	695 <b>9307</b>
Sequence 13972	AF159295	Genbank	571 <b>4635</b>
Sequence 13973	AF161384	Genbank	684 <b>118</b> 1

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Sequence 13974	AF161389	Genbank	6841191
Sequence 13975	AF161428	Genbank	6841269
Sequence 13976	AF161448	Genbank	6841309
Sequence 13977	AF161464	Genbank	6841451
Sequence 13978	AF161479	Genbank	6841481
Sequence 13979	AF161481	Genbank	6841485
Sequence 13980	AF161485	Genbank	
Sequence 13981	AF161500	Genbank	6841493
Sequence 13982	AF161503	Genbank	6841523
Sequence 13983	AF161509	Genbank	6841529
Sequence 13984	AF161512	Genbank	6841541
Sequence 13985	AF161541	Genbank	6841547
Sequence 13986	AF161556		6841349
Sequence 13987	AF165926	Genbank Genbank	6841379
Sequence 13988	AF168956		5817858
Sequence 13989	AF172066	Genbank	5702387
Sequence 13990	AF176315	Genbank	5733811
Sequence 13991		Genbank	5823349
Sequence 13992	AF181896	Genbank	6272683
Sequence 13993	AF187320	Genbank	6164847
	AF187554	Genbank	6653225
Sequence 13994	AF188745	Genbank	6425043
Sequence 13995	AF189712	Genbank	6760474
Sequence 13996	AF190465	Genbank	6120105
Sequence 13997	AF191018	Genbank	6457339
Sequence 13998	AF191298	Genbank	7656642
Sequence 13999	AF201077	Genbank	6456748
Sequence 14000	AF203815	Genbank	6979641
Sequence 14001	AF205588	Genbank	6531675
Sequence 14002	AF207550	Genbank	6470333
Sequence 14003	AF216305	Genbank	6942010
Sequence 14004	AF228703	Genbank	7158868
Sequence 14005	AF228704	Genbank	715 <b>8871</b>
Sequence 14006	AJ000519	Genbank	2739214
Sequence 14007	AJ001014	Genbank	3171909
Sequence 14008	AJ001866	Genbank	2969902
Sequence 14009	AJ001981	Genbank	2546978
Sequence 14010	AJ002030	Genbank	257 <b>0006</b>
Sequence 14011	AJ002308	Genbank	295 <b>9871</b>
Sequence 14012	AJ002385	Genbank	2597930
Sequence 14013	AJ006470	Genbank	3687321
Sequence 14014	AJ010069	Genbank	3483012
Sequence 14015	AJ010597	Genbank	355 <b>9873</b>
Sequence 14016	AJ012409	Genbank	3881975
Sequence 14017	AJ012499	Genbank	5441359
Sequence 14018	AJ223812	Genbank	2894518
Sequence 14019	AJ224442	Genbank	2911586
Sequence 14020	AJ230595	Genbank	4583774
Sequence 14021	AJ230596	Genbank	4583775
Sequence 14022	AJ230605	Genbank	45837 <b>84</b>
Sequence 14023	AJ238243	Genbank	4826530
Sequence 14024	AJ238283	Genbank	4704219
Sequence 14025	AJ238374	Genbank	6634418
Sequence 14026	AJ245719	Genbank	668 <b>8</b> 1 <b>68</b>
Sequence 14027	AJ249731	Genbank	5921469
Sequence 14028	AJ250042	Genbank	6013005
Sequence 14029	AJ250915	Genbank	6996445

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<u>Table 3-5</u>

Sequence 14030	AJ251961	Genbank	6634 <b>45</b> 3
Sequence 14031	AJ271216	Genbank	6966966
Sequence 14032	AKCC0018	Genbank	7019828
Sequence 14033	AKCC0023	Gerbank	7019837
Sequence 14034	AK000035	Gerbank	7019855
Sequence 14035	AK000208	Genbank	7020141
Sequence 14036	AK0C0291	Genbank	7020274
Sequence 14037	AK000348	Genbank	
Sequence 14037	AK000345	Genbank	7020373
Sequence 14039	AK000395		7020402
Sequence 14040	AK000395 AK000426	Genbank	7020455
		Genbank	7020505
Sequence 14041	AK000434	Genbank	7020520
Sequence 14042	AK000453	Genbank	7020550
Sequence 14043	AK000462	Genbank	7020566
Sequence 14044	AK000470	Genbank	7020580
Sequence 14045	AK000507	Genbank	7020644
Sequence 14046	AK000548	Genbank	7020717
Sequence 14047	AK000560	Genbank	7020738
Sequence 14048	AK000571	Genbank	7020756
Sequence 14049	AK000585	Genbank	702 <b>077</b> 9
Sequence 14050	AK000615	Genbank	7020827
Sequence 14051	AK000620	Genbank	7020834
Sequence 14052	AK000624	Genbank	7020839
Sequence 14053	AK000655	Genbank	7020888
Sequence 14054	AK000667	Genbank	7020906
Sequence 14055	AK000669	Genbank	7020909
Sequence 14056	AK000721	Genbank	7020985
Sequence 14057	AK000725	Genbank	7020990
Sequence 14058	AK000739	Genbank	7021014
Sequence 14059	AK000 <b>74</b> 5	Genbank	7021025
Sequence 14060	AK000752	Genbank	7021035
Sequence 14061	AK000765	Genbank	7021058
Sequence 14062	AK000800	Genbank	7021104
Sequence 14063	AK000818	Genbank	7021128
Sequence 14064	AK000826	Genbank	7021139
Sequence 14065	AK000866	Genbank	7021190
Sequence 14066	AK000891	Genbank	7021840
Sequence 14067	AK000933	Genbank	7021908
Sequence 14068	AK000934	Genbank	7021 <b>909</b>
Sequence 14069	AK000959	Genbank	7021945
Sequence 14070	AK000991	Genbank	7021994
Sequence 14071	AK001006	Genbank	7022017
Sequence 14072	AK001093	Genbank	7022142
Sequence 14073	AK001105	Genbank	7022160
Sequence 14074	AK001154	Genbank	7022228
Sequence 14075	AK001237	Genbank	7022366
Sequence 14076	AK001285	Genbank	7022444
Sequence 14077	AK001330	Genbank	7022520
Sequence 14078	AK001387	Genbank	7022615
Sequence 14079	AK001413	Genbank	7022655
Sequence 14080	AK001441	Genbank	7022699
Sequence 14081	AK001451	Genbank	7022717
Sequence 14082	AK001469	Genbank	7022747
Sequence 14083	AK001493	Genbank	7022783
Sequence 14084	AK001536	Genbank	7022851
Sequence 14085	AK001571	Genbank	7022906
-04400106 14003	711001371	Gennank	1022900

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<u>Table 3-5</u>

Sequence 14086         AK001656         Genbank         7023047           Sequence 14087         AK001690         Genbank         7023105           Sequence 14088         AK001691         Genbank         7023105           Sequence 14090         AK001717         Genbank         7023151           Sequence 14091         AK001747         Genbank         7023203           Sequence 14092         AK001762         Genbank         7023231           Sequence 14093         AK001875         Genbank         7023382           Sequence 14094         AK001877         Genbank         7023416           Sequence 14095         AK001878         Genbank         7023418           Sequence 14096         AK001878         Genbank         7023418           Sequence 14097         AK002002         Genbank         7023620           Sequence 14099         AK002046         Genbank         7023781           Sequence 14101         AK002169         Genbank         7023781           Sequence 14101         AK002190         Genbank         7023984           Sequence 14101         AL008582         Genbank         7023914           Sequence 14102         AL008721         Genbank         2793914				
Sequence 14087         AK001690         Genbank         7023105           Sequence 14088         AK001691         Genbank         7023105           Sequence 14089         AK001717         Genbank         7023153           Sequence 14090         AK001747         Genbank         7023153           Sequence 14091         AK001762         Genbank         7023203           Sequence 14093         AK001855         Genbank         7023416           Sequence 14094         AK001877         Genbank         7023418           Sequence 14095         AK001878         Genbank         7023418           Sequence 14096         AK001961         Genbank         7023549           Sequence 14097         AK002002         Genbank         7023620           Sequence 14099         AK002102         Genbank         7023781           Sequence 14100         AK002190         Genbank         7023781           Sequence 14101         AK002190         Genbank         7023781           Sequence 14102         AL008582         Genbank         7023914           Sequence 14103         AL008582         Genbank         5360981           Sequence 14106         AL008525         Genbank         5360981	Sequence 1408	6 AK001656	Genbank	7023047
Sequence 14088         AK001691         Genbank         7023100           Sequence 14089         AK001717         Genbank         7023153           Sequence 14090         AK001747         Genbank         7023203           Sequence 14091         AK001747         Genbank         7023203           Sequence 14092         AK001875         Genbank         7023382           Sequence 14094         AK001877         Genbank         7023418           Sequence 14095         AK001878         Genbank         7023418           Sequence 14097         AK002002         Genbank         7023549           Sequence 14098         AK002002         Genbank         7023620           Sequence 14099         AK002012         Genbank         7023781           Sequence 14100         AK002169         Genbank         7023884           Sequence 14101         AK002190         Genbank         7023914           Sequence 14102         AL008582         Genbank         7023914           Sequence 14103         AL008582         Genbank         5360981           Sequence 14104         AL008721         Genbank         6599068           Sequence 14105         AL008725         Genbank         2591595				
Sequence 14089         AK001717         Genbank         7023151           Sequence 14090         AK001747         Genbank         7023153           Sequence 14091         AK001747         Genbank         7023231           Sequence 14093         AK001855         Genbank         7023231           Sequence 14094         AK001877         Genbank         7023418           Sequence 14095         AK001878         Genbank         7023418           Sequence 14096         AK001961         Genbank         7023549           Sequence 14097         AK002002         Genbank         7023549           Sequence 14099         AK002102         Genbank         7023781           Sequence 14099         AK002102         Genbank         7023781           Sequence 14101         AK002190         Genbank         7023781           Sequence 14102         AL008582         Genbank         7023914           Sequence 14103         AL008582         Genbank         5360981           Sequence 14103         AL008582         Genbank         5360981           Sequence 14105         AL008725         Genbank         259155           Sequence 14106         AL008725         Genbank         259155				
Sequence 14090         AK001718         Genbank         7023153           Sequence 14091         AK001762         Genbank         7023203           Sequence 14093         AK001762         Genbank         7023231           Sequence 14094         AK001875         Genbank         7023418           Sequence 14096         AK001878         Genbank         7023418           Sequence 14097         AK002002         Genbank         7023549           Sequence 14099         AK002002         Genbank         7023620           Sequence 14099         AK0020162         Genbank         7023689           Sequence 14099         AK002169         Genbank         7023781           Sequence 14100         AK002169         Genbank         7023781           Sequence 14100         AK002169         Genbank         7023781           Sequence 14101         AK002169         Genbank         7023781           Sequence 14102         AL008522         Genbank         7023781           Sequence 14103         AL008521         Genbank         659068           Sequence 14104         AL008721         Genbank         299155           Sequence 14105         AL008721         Genbank         2995195				
Sequence 14091         AK001747         Genbank         7023203           Sequence 14092         AK001762         Genbank         7023231           Sequence 14094         AK001875         Genbank         70233416           Sequence 14095         AK001878         Genbank         7023418           Sequence 14096         AK001961         Genbank         7023549           Sequence 14097         AK002002         Genbank         7023689           Sequence 14099         AK002046         Genbank         7023689           Sequence 14100         AK002102         Genbank         7023689           Sequence 14100         AK002102         Genbank         7023781           Sequence 14100         AK002190         Genbank         7023884           Sequence 14101         AK002190         Genbank         7023914           Sequence 14102         AL008582         Genbank         6599068           Sequence 14103         AL008583         Genbank         6599068           Sequence 14105         AL008725         Genbank         2791551           Sequence 14106         AL009951         Genbank         2995195           Sequence 14109         AL020995         Genbank         2853183				
Sequence 14092         AK001762         Genbank         7023231           Sequence 14093         AK001855         Genbank         7023382           Sequence 14095         AK001877         Genbank         7023418           Sequence 14096         AK001961         Genbank         7023418           Sequence 14097         AK002002         Genbank         7023620           Sequence 14098         AK002046         Genbank         7023689           Sequence 14099         AK002102         Genbank         7023781           Sequence 14100         AK002190         Genbank         7023781           Sequence 14101         AK002190         Genbank         7023781           Sequence 14101         AK002190         Genbank         7023781           Sequence 14101         AK008582         Genbank         7023914           Sequence 14103         AL008582         Genbank         5360981           Sequence 14104         AL008721         Genbank         2590988           Sequence 14105         AL008721         Genbank         2995195           Sequence 14107         AL009711         Genbank         2995195           Sequence 14108         AL020993         Genbank         2853173				
Sequence 14093         AK001855         Genbank         7023382           Sequence 14094         AK001878         Genbank         7023418           Sequence 14096         AK001961         Genbank         7023549           Sequence 14097         AK002002         Genbank         7023620           Sequence 14099         AK0020102         Genbank         7023689           Sequence 14099         AK002102         Genbank         7023781           Sequence 14100         AK002169         Genbank         7023781           Sequence 14101         AK002169         Genbank         7023781           Sequence 14102         AL008582         Genbank         7023914           Sequence 14103         AL008592         Genbank         569968           Sequence 14104         AL008725         Genbank         6599068           Sequence 14105         AL008725         Genbank         2791551           Sequence 14106         AL009725         Genbank         2791551           Sequence 14107         AL009781         Genbank         2853179           Sequence 14108         AL020993         Genbank         2853183           Sequence 14110         AL021368         Genbank         2853183				
Sequence 14094         AK001877         Genbank         7023416           Sequence 14095         AK001878         Genbank         7023418           Sequence 14096         AK001961         Genbank         7023549           Sequence 14097         AK002002         Genbank         7023620           Sequence 14099         AK002046         Genbank         7023689           Sequence 14100         AK002169         Genbank         7023781           Sequence 14101         AK002190         Genbank         7023914           Sequence 14102         AL008582         Genbank         5360981           Sequence 14103         AL008639         Genbank         5360981           Sequence 14105         AL008725         Genbank         6599068           Sequence 14106         AL009725         Genbank         2791551           Sequence 14107         AL009781         Genbank         2853179           Sequence 14108         AL020993         Genbank         2853183           Sequence 14110         AL021069         Genbank         2853183           Sequence 14111         AL021069         Genbank         286890           Sequence 14111         AL021920         Genbank         2826890				
Sequence 14095         AK001878         Genbank         7023418           Sequence 14096         AK001961         Genbank         7023549           Sequence 14097         AK002002         Genbank         7023620           Sequence 14099         AK002046         Genbank         7023683           Sequence 14100         AK002169         Genbank         7023781           Sequence 14101         AK002190         Genbank         7023914           Sequence 14102         AL008582         Genbank         5360981           Sequence 14103         AL008539         Genbank         6599068           Sequence 14104         AL008721         Genbank         6599068           Sequence 14105         AL003725         Genbank         2995195           Sequence 14106         AL003725         Genbank         2995195           Sequence 14107         AL009181         Genbank         2985195           Sequence 14108         AL020993         Genbank         2853179           Sequence 14110         AL021069         Genbank         2853183           Sequence 14111         AL021669         Genbank         2863183           Sequence 14111         AL021669         Genbank         2826890	•			
Sequence 14096         AK001961         Genbank         7023549           Sequence 14097         AK002002         Genbank         7023620           Sequence 14098         AK002046         Genbank         7023689           Sequence 14099         AK002102         Genbank         7023781           Sequence 14100         AK002199         Genbank         7023884           Sequence 14101         AK002199         Genbank         7023914           Sequence 14102         AL008582         Genbank         5360981           Sequence 14103         AL008529         Genbank         6599068           Sequence 14105         AL008721         Genbank         2791551           Sequence 14106         AL009725         Genbank         2791551           Sequence 14107         AL009181         Genbank         2853179           Sequence 14108         AL020993         Genbank         380107           Sequence 14110         AL021069         Genbank         2853183           Sequence 14111         AL021969         Genbank         2853183           Sequence 14111         AL021969         Genbank         2864535           Sequence 14111         AL021546         Genbank         2864535				
Sequence 14097         AK002002         Genbank         7023620           Sequence 14098         AK002046         Genbank         7023689           Sequence 14109         AK002102         Genbank         7023781           Sequence 14101         AK002169         Genbank         7023814           Sequence 14102         AL008582         Genbank         5360981           Sequence 14103         AL008639         Genbank         5569968           Sequence 14104         AL008721         Genbank         2791551           Sequence 14105         AL008725         Genbank         2791551           Sequence 14107         AL009181         Genbank         2995195           Sequence 14108         AL009951         Genbank         2995195           Sequence 14109         AL020993         Genbank         2853179           Sequence 14110         AL021069         Genbank         2853183           Sequence 14111         AL021546         Genbank         2853183           Sequence 14111         AL021546         Genbank         3281985           Sequence 14113         AL022165         Genbank         3281985           Sequence 14114         AL022165         Genbank         426451				
Sequence 14098         AK002046         Genbank         7023699           Sequence 14099         AK002102         Genbank         7023781           Sequence 14100         AK002199         Genbank         7023844           Sequence 14101         AK002190         Genbank         7023914           Sequence 14102         AL008582         Genbank         5360981           Sequence 14104         AL008639         Genbank         6599068           Sequence 14105         AL008725         Genbank         2791551           Sequence 14106         AL009725         Genbank         2995195           Sequence 14107         AL009181         Genbank         2995195           Sequence 14109         AL020993         Genbank         2853179           Sequence 14110         AL021069         Genbank         2853183           Sequence 14111         AL02168         Genbank         2853183           Sequence 14111         AL021546         Genbank         286893           Sequence 14111         AL021546         Genbank         3281985           Sequence 14114         AL022238         Genbank         426451           Sequence 14115         AL022238         Genbank         426471				
Sequence 14109         AK002102         Genbank         7023781           Sequence 14101         AK002169         Genbank         7023884           Sequence 14101         AK002190         Genbank         7023914           Sequence 14102         AL008582         Genbank         5360981           Sequence 14103         AL008639         Genbank         559068           Sequence 14104         AL008721         Genbank         3171883           Sequence 14105         AL008725         Genbank         2791551           Sequence 14106         AL009051         Genbank         2995195           Sequence 14107         AL009181         Genbank         2853179           Sequence 14109         AL020993         Genbank         2853183           Sequence 14110         AL021368         Genbank         2853183           Sequence 14111         AL021368         Genbank         286890           Sequence 14113         AL021960         Genbank         2826890           Sequence 14114         AL022165         Genbank         3281985           Sequence 14115         AL022166         Genbank         4264535           Sequence 14116         AL022238         Genbank         426471				
Sequence 14100         AK002169         Genbank         7023884           Sequence 14101         AK002190         Genbank         7023914           Sequence 14102         AL008582         Genbank         5360981           Sequence 14103         AL008539         Genbank         6599068           Sequence 14104         AL008721         Genbank         2791551           Sequence 14105         AL008725         Genbank         2995195           Sequence 14107         AL009181         Genbank         2995195           Sequence 14108         AL020993         Genbank         2985195           Sequence 14109         AL020995         Genbank         3980107           Sequence 14110         AL021069         Genbank         2853183           Sequence 14111         AL021368         Genbank         2853183           Sequence 14111         AL021368         Genbank         286890           Sequence 14111         AL021920         Genbank         3264535           Sequence 14111         AL022238         Genbank         3264535           Sequence 14115         AL022238         Genbank         4826471           Sequence 14116         AL0222302         Genbank         4826471	•			
Sequence 14101         AK002190         Genbank         7023914           Sequence 14102         AL008582         Genbank         5360981           Sequence 14103         AL008639         Genbank         6599068           Sequence 14104         AL008721         Genbank         2791551           Sequence 14105         AL008725         Genbank         2995195           Sequence 14106         AL009051         Genbank         2995195           Sequence 14107         AL009181         Genbank         2985179           Sequence 14109         AL020993         Genbank         2985179           Sequence 14110         AL021069         Genbank         2853183           Sequence 14111         AL021368         Genbank         2853183           Sequence 14111         AL021466         Genbank         286890           Sequence 14111         AL021920         Genbank         3281985           Sequence 14111         AL022238         Genbank         3281985           Sequence 14111         AL022238         Genbank         4176442           Sequence 14116         AL0222302         Genbank         426471           Sequence 14117         AL022302         Genbank         420326				
Sequence 14102         AL008582         Genbank         5360981           Sequence 14103         AL008639         Genbank         6599068           Sequence 14104         AL008721         Genbank         3171883           Sequence 14106         AL008725         Genbank         2791551           Sequence 14107         AL009181         Genbank         2995195           Sequence 14108         AL020993         Genbank         3980107           Sequence 14109         AL020995         Genbank         2881341           Sequence 14110         AL021069         Genbank         2853183           Sequence 14111         AL021368         Genbank         2826890           Sequence 14112         AL021546         Genbank         3264535           Sequence 14113         AL021920         Genbank         3264535           Sequence 14114         AL022165         Genbank         3264535           Sequence 14115         AL022238         Genbank         426471           Sequence 14116         AL0222312         Genbank         426471           Sequence 14118         AL022312         Genbank         4200326           Sequence 14119         AL022313         Genbank         4200326	•		Genbank	7023884
Sequence 14103         AL008639         Genbank         6599068           Sequence 14104         AL008721         Genbank         3171883           Sequence 14105         AL008725         Genbank         2791551           Sequence 14106         AL009051         Genbank         2995195           Sequence 14107         AL009181         Genbank         2853179           Sequence 14109         AL020993         Genbank         3980107           Sequence 14110         AL021069         Genbank         2853183           Sequence 14111         AL021368         Genbank         3080468           Sequence 14111         AL021546         Genbank         2826890           Sequence 14112         AL021546         Genbank         3284535           Sequence 14113         AL021920         Genbank         3281985           Sequence 14114         AL022365         Genbank         4176442           Sequence 14115         AL022388         Genbank         4176442           Sequence 14116         AL022302         Genbank         420326           Sequence 14117         AL022302         Genbank         4200326           Sequence 14119         AL022313         Genbank         4200326	•		Genbank	
Sequence 14104         AL008725         Genbank         2791551           Sequence 14106         AL008725         Genbank         2791551           Sequence 14107         AL009181         Genbank         2995195           Sequence 14108         AL020993         Genbank         2853179           Sequence 14109         AL020995         Genbank         3880107           Sequence 14110         AL021069         Genbank         2853183           Sequence 14111         AL021368         Genbank         2826890           Sequence 14112         AL021546         Genbank         2826890           Sequence 14113         AL021920         Genbank         3284535           Sequence 14114         AL022165         Genbank         3281985           Sequence 14115         AL022238         Genbank         4176442           Sequence 14117         AL022302         Genbank         4826471           Sequence 14118         AL022312         Genbank         4200326           Sequence 14119         AL022313         Genbank         4200326           Sequence 14120         AL022313         Genbank         4914501           Sequence 14121         AL022320         Genbank         4914513	•		Genbank	5360981
Sequence 14105         AL008725         Genbank         2791551           Sequence 14106         AL009051         Genbank         2995195           Sequence 14107         AL009181         Genbank         2853179           Sequence 14108         AL020993         Genbank         3980107           Sequence 14110         AL021069         Genbank         2853183           Sequence 14111         AL021368         Genbank         2853183           Sequence 14112         AL021546         Genbank         2826890           Sequence 14113         AL021920         Genbank         3264535           Sequence 14114         AL022165         Genbank         3281985           Sequence 14115         AL022238         Genbank         4826471           Sequence 14116         AL0222302         Genbank         4826471           Sequence 14118         AL022312         Genbank         4914501           Sequence 14119         AL022313         Genbank         490326           Sequence 14110         AL022313         Genbank         4914501           Sequence 14120         AL022313         Genbank         4914501           Sequence 14121         AL022328         Genbank         5263010			Genbank	6599068
Sequence 14106         AL009051         Genbank         2995195           Sequence 14107         AL009181         Genbank         2853179           Sequence 14108         AL020993         Genbank         3980107           Sequence 14110         AL021069         Genbank         5881341           Sequence 14111         AL021368         Genbank         2853183           Sequence 14111         AL021546         Genbank         2826890           Sequence 14113         AL021920         Genbank         3281985           Sequence 14114         AL021655         Genbank         3281985           Sequence 14115         AL022238         Genbank         4176442           Sequence 14116         AL022238         Genbank         426471           Sequence 14117         AL022302         Genbank         420442           Sequence 14119         AL022312         Genbank         4200326           Sequence 14119         AL022313         Genbank         4200326           Sequence 14120         AL022316         Genbank         4200326           Sequence 14121         AL022328         Genbank         5263010           Sequence 14123         AL022324         Genbank         5263010	Sequence 1410	4 AL008721	Genbank	3171883
Sequence 14107         AL009181         Genbank         2853179           Sequence 14108         AL020993         Genbank         3980107           Sequence 14110         AL020995         Genbank         5881341           Sequence 14111         AL021069         Genbank         2853183           Sequence 14111         AL021546         Genbank         2826890           Sequence 14113         AL021920         Genbank         3284535           Sequence 14114         AL022165         Genbank         3281985           Sequence 14115         AL022238         Genbank         4176442           Sequence 14116         AL022238         Genbank         4826471           Sequence 14117         AL022302         Genbank         4826471           Sequence 14117         AL022302         Genbank         4914501           Sequence 14119         AL022313         Genbank         4914501           Sequence 14120         AL022316         Genbank         4200326           Sequence 14121         AL022320         Genbank         5263010           Sequence 14122         AL022328         Genbank         5263010           Sequence 14123         AL022394         Genbank         5679748	Sequence 14109	5 AL008725	Genbank	2791551
Sequence 14108         AL020993         Genbank         3980107           Sequence 14109         AL020995         Genbank         5881341           Sequence 14110         AL021069         Genbank         2853183           Sequence 14111         AL021368         Genbank         3080468           Sequence 14111         AL021546         Genbank         2826890           Sequence 14113         AL021920         Genbank         3264535           Sequence 14114         AL022238         Genbank         3281985           Sequence 14115         AL022238         Genbank         4176442           Sequence 14116         AL022238         Genbank         426471           Sequence 14117         AL022302         Genbank         420326           Sequence 14118         AL022312         Genbank         4200326           Sequence 14119         AL022316         Genbank         4691242           Sequence 14120         AL022316         Genbank         420326           Sequence 14121         AL022328         Genbank         5263010           Sequence 14123         AL022394         Genbank         5263010           Sequence 14124         AL022395         Genbank         535875	Sequence 14100	6 AL009051	Genbank	2995195
Sequence 14109         AL020995         Genbank         5881341           Sequence 14110         AL021069         Genbank         2853183           Sequence 14111         AL021368         Genbank         3080468           Sequence 14112         AL021546         Genbank         3264535           Sequence 14113         AL021920         Genbank         3264535           Sequence 14114         AL0222165         Genbank         3281985           Sequence 14115         AL022238         Genbank         4176442           Sequence 14116         AL0222302         Genbank         4826471           Sequence 14117         AL022302         Genbank         4914501           Sequence 14118         AL022312         Genbank         4914501           Sequence 14119         AL022313         Genbank         4200326           Sequence 14120         AL022316         Genbank         4691242           Sequence 14121         AL022320         Genbank         4691242           Sequence 14122         AL022328         Genbank         5263010           Sequence 14123         AL022394         Genbank         5263010           Sequence 14124         AL022394         Genbank         567948	Sequence 14107	7 AL009181	Genbank	2853179
Sequence 14109         AL020995         Genbank         5881341           Sequence 14110         AL021069         Genbank         2853183           Sequence 14111         AL021368         Genbank         3080468           Sequence 14112         AL021546         Genbank         3264535           Sequence 14113         AL021920         Genbank         3264535           Sequence 14114         AL0222165         Genbank         3281985           Sequence 14115         AL022238         Genbank         4176442           Sequence 14116         AL0222302         Genbank         4826471           Sequence 14117         AL022302         Genbank         4914501           Sequence 14118         AL022312         Genbank         4914501           Sequence 14119         AL022313         Genbank         4200326           Sequence 14120         AL022316         Genbank         4691242           Sequence 14121         AL022320         Genbank         4691242           Sequence 14122         AL022328         Genbank         5263010           Sequence 14123         AL022394         Genbank         5263010           Sequence 14124         AL022394         Genbank         567948	Sequence 1410	3 AL020993	Genbank	3980107
Sequence 14110         AL021069         Genbank         2853183           Sequence 14111         AL021368         Genbank         3080468           Sequence 14112         AL021546         Genbank         2826890           Sequence 14113         AL021920         Genbank         3264535           Sequence 14114         AL022165         Genbank         3281985           Sequence 14115         AL022238         Genbank         4176442           Sequence 14116         AL0222302         Genbank         4826471           Sequence 14117         AL022302         Genbank         4914501           Sequence 14118         AL022312         Genbank         4914501           Sequence 14119         AL022313         Genbank         4200326           Sequence 14120         AL022316         Genbank         4691242           Sequence 14121         AL022320         Genbank         4914513           Sequence 14122         AL022328         Genbank         5263010           Sequence 14123         AL022394         Genbank         5679748           Sequence 14125         AL022325         Genbank         5679748           Sequence 14127         AL023494         Genbank         3355875	Sequence 14109	AL020995	Genbank	
Sequence 14111         AL021368         Genbank         3080468           Sequence 14112         AL021546         Genbank         2826890           Sequence 14113         AL021920         Genbank         3264535           Sequence 14114         AL022165         Genbank         3281985           Sequence 14115         AL022238         Genbank         4176442           Sequence 14116         AL0222302         Genbank         4826471           Sequence 14117         AL022302         Genbank         4914501           Sequence 14118         AL022312         Genbank         4914501           Sequence 14119         AL022313         Genbank         4914501           Sequence 14120         AL022316         Genbank         4691242           Sequence 14121         AL022320         Genbank         4914513           Sequence 14122         AL022328         Genbank         5263010           Sequence 14123         AL022394         Genbank         581758           Sequence 14124         AL022395         Genbank         3355875           Sequence 14125         AL023284         Genbank         3355875           Sequence 14128         AL023494         Genbank         3350031	Sequence 14110	AL021069	Genbank	
Sequence 14112         AL021546         Genbank         2826890           Sequence 14113         AL021920         Genbank         3264535           Sequence 14114         AL022165         Genbank         3281985           Sequence 14115         AL022238         Genbank         4176442           Sequence 14116         AL022240         Genbank         4826471           Sequence 14117         AL022302         Genbank         5262844           Sequence 14118         AL022312         Genbank         4914501           Sequence 14119         AL022313         Genbank         4200326           Sequence 14120         AL022316         Genbank         4601242           Sequence 14121         AL022320         Genbank         4914513           Sequence 14122         AL022328         Genbank         5263010           Sequence 14123         AL022394         Genbank         5263010           Sequence 14124         AL022394         Genbank         4468287           Sequence 14125         AL022395         Genbank         5679748           Sequence 14126         AL023284         Genbank         5360992           Sequence 14128         AL023494         Genbank         3550031				
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Sequence 14114         AL022165         Genbank         3281985           Sequence 14115         AL022238         Genbank         4176442           Sequence 14116         AL022240         Genbank         4826471           Sequence 14117         AL022302         Genbank         5262844           Sequence 14118         AL022312         Genbank         4914501           Sequence 14119         AL022313         Genbank         4200326           Sequence 14120         AL022316         Genbank         4691242           Sequence 14121         AL022320         Genbank         4914513           Sequence 14122         AL022328         Genbank         5263010           Sequence 14123         AL022394         Genbank         9581758           Sequence 14124         AL022395         Genbank         4468287           Sequence 14125         AL022725         Genbank         5355875           Sequence 14126         AL023284         Genbank         3355875           Sequence 14127         AL023494         Genbank         3550031           Sequence 14128         AL023798         Genbank         3288439           Sequence 14130         AL031003         Genbank         3688349				
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Sequence 14133         AL031274         Genbank         3421083           Sequence 14134         AL031287         Genbank         4914533           Sequence 14135         AL031295         Genbank         4376011           Sequence 14136         AL031368         Genbank         4455593           Sequence 14137         AL031390         Genbank         5002609           Sequence 14138         AL031428         Genbank         5931719           Sequence 14139         AL031431         Genbank         4938290           Sequence 14140         AL031432         Genbank         4375969		_		
Sequence 14134         AL031287         Genbank         4914533           Sequence 14135         AL031295         Genbank         4376011           Sequence 14136         AL031368         Genbank         4455593           Sequence 14137         AL031390         Genbank         5002609           Sequence 14138         AL031428         Genbank         5931719           Sequence 14139         AL031431         Genbank         4938290           Sequence 14140         AL031432         Genbank         4375969				
Sequence 14135         AL031295         Genbank         4376011           Sequence 14136         AL031368         Genbank         4455593           Sequence 14137         AL031390         Genbank         5002609           Sequence 14138         AL031428         Genbank         5931719           Sequence 14139         AL031431         Genbank         4938290           Sequence 14140         AL031432         Genbank         4375969				
Sequence 14136         AL031368         Genbank         4455593           Sequence 14137         AL031390         Genbank         5002609           Sequence 14138         AL031428         Genbank         5931719           Sequence 14139         AL031431         Genbank         4938290           Sequence 14140         AL031432         Genbank         4375969				
Sequence 14137         AL031390         Genbank         5002609           Sequence 14138         AL031428         Genbank         5931719           Sequence 14139         AL031431         Genbank         4938290           Sequence 14140         AL031432         Genbank         4375969				
Sequence 14138         AL031428         Genbank         5931719           Sequence 14139         AL031431         Genbank         4938290           Sequence 14140         AL031432         Genbank         4375969	•			
Sequence 14139         AL031431         Genbank         4938290           Sequence 14140         AL031432         Genbank         4375969				
Sequence 14140 AL031432 Genbank 4375969				
				4938290
Sequence 14141 AL031433 Genbank 4826442			Genbank	4375969
	Sequence 14141	AL031433	Genbank	4826442

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<u>Table 3-5</u>

Sequence 14142	AL031447	Genbank	4826431
Sequence 14143	AL031597	Genbank	4775640
Sequence 14144	AL031662	Genbank	9716901
Sequence 14145	AL031667	Genbank	9408727
Sequence 14146	AL031681	Genbank	9408729
Sequence 14147	AL031710	Genbank	9798435
Sequence 14148	AL031716	Genbank	4826474
Sequence 14149	AL031732	Genbank	4581343
Sequence 14150	AL031781	Genbank	4038570
Sequence 14151	AL032821	Genbank	4572584
Sequence 14152	AL033377	Genbank	4826462
Secuence 14153	AL033397	Genbank	4902626
Sequence 14154	AL033522	Genbank	4191258
Sequence 14155	AL033531	Genbank	
Sequence 14156	AL033543	Genbank	6807582 4581346
Sequence 14157	AL034417	Genbank	
Sequence 14158	AL034548	Genbank	5102616
Sequence 14159	AL035209	Genbank	7263904
Sequence 14160	AL035413	Genbank	4160217
Sequence 14161	AL035420	Genbank	6010110
Sequence 14162	AL035425	Genbank	5918161
Sequence 14163	AL035458	Genbank	6465823
Sequence 14164	AL035541	Genbank	6624641
Sequence 14165	AL035652	Genbank	9581764
Sequence 14166	AL035681	Genbank	9650702
Sequence 14167	AL035699	Genbank	4902689
Sequence 14168	AL049249	Genbank	4826515
Sequence 14169	AL049282	Genbank	4499987
Sequence 14170	AL049404	Genbank	4500041
Sequence 14171	AL049557	Genbank	4500192 5506765
Sequence 14172	AL049563	Genbank	5596765
Sequence 14173	AL049564	Genbank	4902753 4902757
Sequence 14174	AL049589	Genbank	5679567
Sequence 14175	AL049696	Genbank	5931893
Sequence 14176	AL049742	Genbank	5419783
Sequence 14177	AL049761	Genbank	5738437
Sequence 14178	AL049766	Genbank	5763746
Sequence 14179	AL049776	Genbank	6681704
Sequence 14180	AL049779	Genbank	8176894
Sequence 14181	AL049829	Genbank	8217859
Sequence 14182	AL049844	Genbank	5924019
Sequence 14183	AL049869	Genbank	8217861
Sequence 14184	AL049929	Genbank	4884174
Sequence 14185	AL049968	Genbank	4884217
Sequence 14186	AL049987	Genbank	4884238
Sequence 14187	AL050021	Genbank	4884264
Sequence 14188	AL050027	Genbank	4884266
Sequence 14189	AL050035	Genbank	4884276
Sequence 14190	AL050083	Genbank	4884317
Sequence 14191	AL050161	Genbank	4884375
Sequence 14192	AL050170	Genbank	4884165
Sequence 14193	AL050215	Genbank	4884454
Sequence 14194	AL050273	Genbank	4886450
Sequence 14195	AL050275	Genbank	4886500
Sequence 14196	AL050290	Genbank	4886512
Sequence 14197	ALC50343	Genbank	6137008
			5.57500

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**Table 3-5** 

Sequence 14198	AL050363	Genbank	4914597
Sequence 14199	AL050383	Genbank	4914585
Sequence 14200	AL050392	Genbank	4914613
Sequence 14201	AL078591	Genbank	6562085
Sequence 14202	AL078621	Genbank	6013067
Sequence 14203	AL079295	Genbank	5102607
Sequence 14204	AL079304	Genbank	8176897
Sequence 14205	AL079342	Genbank	6018784
Sequence 14206	AL080088	Genbank	5262504
Sequence 14207	AL080089	Genbank	526250 <b>6</b>
Sequence 14208	AL080097	Genbank	5262519
Sequence 14209	AL080109	Genbank	5262535
Sequence 14210	AL080135	Genbank	5262576
Sequence 14211	AL080144	Genbank	5262592
Sequence 14212	AL080186	Genbank	5262664
Sequence 14213	AL080241	Genbank	5730204
Sequence 14214	AL080250	Genbank	5804917
Sequence 14215	AL080276	Genbank	5763753
Sequence 14216	AL096773	Genbank	5918011
Sequence 14217	AL096803	Genbank	6010162
Sequence 14218	AL109657	Genbank	6136991
Sequence 14219	AL109684	Genbank	5689805
Sequence 14220	AL109925	Genbank	7105935
Sequence 14221	AL110183	Genbank	5817095
Sequence 14222	AL110185	Genbank	
Sequence 14223	AL110193	Genbank	5817098 5817109
Sequence 14224	AL110194	Genbank	5817111
Sequence 14225	AL110202	Genbank	5817121
Sequence 14226	AL110202	Genbank	5817127
Sequence 14227	AL110262	Genbank	5817224
Sequence 14228	AL117334	Genbank	7248316
Sequence 14229	AL117424	Genbank	5911856
Sequence 14230	AL117430	Genbank	5911865
Sequence 14231	AL117442	Genbank	5911885
Sequence 14232	AL117558	Genbank	5912097
Sequence 14233	AL117599	Genbank	5912167
Sequence 14234	AL117621	Genbank	5912202
Sequence 14235	AL117648	Genbank	5912240
Sequence 14236	AL117662	Genbank	5912257
Sequence 14237	AL117666	Genbank	5912264
Sequence 14238	AL121576	Genbank	8176910
Sequence 14239	AL121580	Genbank	6456854
Sequence 14240	AL121603	Genbank	6434634
Sequence 14241	AL121656	Genbank	7159617
Sequence 14242	AL121782	Genbank	6456831
Sequence 14243	AL121790	Genbank	8919824
Sequence 14244	AL121913	Genbank	7161781
Sequence 14245	AL121963	Genbank	7161783
Sequence 14246	AL122003	Genbank	6580480
Sequence 14247	AL122013	Genbank	7242500
Sequence 14248	AL122013	Genbank	6093244
Sequence 14249	AL122066	Genbank	6102860
Sequence 14250	AL122089	Genbank	6102899
Sequence 14251	AL133051	Genbank	6453468
Sequence 14252	AL133060	Genbank	6453483
Sequence 14253	AL133094	Genbank	6453548
234401100 17200	/ IE 100007	Genbank	0-000-0

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# <u>Table 3-5</u>

Sequence 14254	AL133238	Genbank	6624577
Sequence 14255	AL133241	Genbank	7630038
Sequence 14256	AL133243	Genbank	6491713
Sequence 14257	AL133246	Genbank	-
Sequence 14258	AL133370	Genbank	7594587
Sequence 14259	AL133453	_	8248722
Sequence 14260		Genbank	7009594
	AL133455	Genbank	7019734
Sequence 14261	AL133555	Genbank	6599122
Sequence 14262	AL135744	Genbank	6 <b>682291</b>
Sequence 14263	AL135858	Genbank	7159622
Sequence 14264	AL136296	Genbank	8217913
Sequence 14265	AL136504	Genbank	7018294
Sequence 14266	AL137440	Genbank	68 <b>08001</b>
Sequence 14267	AL137450	Genbank	6808024
Sequence 14268	AL137489	Genbank	6808110
Sequence 14269	AL137496	Genbank	6808137
Sequence 14270	AL137553	Genbank	6808238
Sequence 14271	AL137567	Genbank	6808276
Sequence 14272	AL137659	Genbank	6807778
Sequence 14273	AL137673	Genbank	6807841
Sequence 14274	AL137681	Genbank	
Sequence 14275	AL137714	Genbank	6807931
Sequence 14276	AL137750	Genbank	6808128
Sequence 14277	AL157730 AL157428	Genbank	6808386
Sequence 14278	AL157464		7018458
Sequence 14279	AP000008	Genbank	7018479
Sequence 14280	AP000031	Genbank	3132318
Sequence 14281	AP000038	Ger.bank Genbank	3132341
Sequence 14282	AP000351		3132348
Sequence 14283	AP000354	Genbank	6069325
Sequence 14284	AP000473	Genbank	5103017
Sequence 14285	AP000525	Genbank Genbank	6942329
Sequence 14286	AP000526		5931503
Sequence 14287	AP000962	Genbank	5931504
Sequence 14288	AP001055	Genbank	6942330
Sequence 14289	AP001035	Genbank	6693605
Sequence 14290	AP001137	Genbank	6970360
Sequence 14291	AP001172	Genbank	6970361
Sequence 14292	AP001172 AP001343	Genbank	6983884
Sequence 14293		Genbank	7209831
_ `	AP001346	Genbank	7209834
Sequence 14294 Sequence 14295	D10493	Genbank	219932
Sequence 14296	D12485	Genbank	219943
Sequence 14297	D14082	Genbank	468933
	D14533	Genbank	286028
Sequence 14298	D14540	Genbank	467511
Sequence 14299	D14658	Genbank	285940
Sequence 14300	D14665	Genbank	6 <b>630617</b>
Sequence 14301	D14696	Genbank	2 <b>85962</b>
Sequence 14302	D14710	Genbank	5 <b>59324</b>
Sequence 14303	D21089	Genbank	475 <b>156</b>
Sequence 14304	D21235	Genbank	498145
Sequence 14305	D21262	Genbank	434764
Sequence 14306	D25215	Genbank	517114
Sequence 14307	D25274	Genbank	464185
Sequence 14308	D26600	Genbank	5 <b>65650</b>
Sequence 14309	D28468	Genbank	460704

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<u>Table 3-5</u>

Sequence 14310	D29011	Genbank	55 <b>852</b> 5
Sequence 14311	D29643	Genbank	473936
Sequence 14312	D29805	Genbank	474986
Sequence 14313	D29954	Genbank	473940
Sequence 14314	D29958	Genbank	
Sequence 14315		Genbank	473948
•	D30648		506337
Sequence 14316	D31885	Genbank	505097
Sequence 14317	D32053	Genbank	2366751
Sequence 14318	· D38112	. Genbank	644480
Sequence 14319	D38524	Genbank	633070
Sequence 14320	D42041	Genbank	5 <b>77294</b>
Sequence 14321	D42063	Genbank	924266
Sequence 14322	D49489	Genbank	1136742
Sequence 14323	D50420	Genbank	2618577
Sequence 14324	D50525	Genbank	1167502
Sequence 14325	D50916	Genbank	1469174
Sequence 14326	D50929	Genbank	1469200
Sequence 14327	D55673	Genbank	870746
Sequence 14328	D63486	Genbank	1469885
Sequence 14329	D63880	Genbank	
		Genbank	961451
Sequence 14330	D64015		2281005
Sequence 14331	D79205	Genbank	1754620
Sequence 14332	D79986	Genbank	1136389
Sequence 14333	D80000	Genbank	1136415
Sequence 14334	D83004	Genbank	1181557
Sequence 14335	D84273	Genbank	3894217
Sequence 14336	D86962	Genbank	1503997
Sequence 14337	D86985	Genbank	6634002
Sequence 14338	D87003	Genbank	2114242
Sequence 14339	D87127	Genbank	1817551
Sequence 14340	D87328	Genbank	1813423
Sequence 14341	D87666	Genbank	1620 <b>016</b>
Sequence 14342	D87684	Genbank	1663703
Sequence 14343	D90209	Genbank	220087
Sequence 14344	E00882	Genbank	2169143
Sequence 14345	E01094	Genbank	2169353
Sequence 14346	E01533	Genbank	2169789
Sequence 14347	E01932	Genbank	2170180
Sequence 14348	E01954	Genbank	2170202
Sequence 14349	E01979	Genbank	2170227
Sequence 14350	E02164	Genbank	2170402
Sequence 14351	E02628	Genbank	2170856
Sequence 14352	E05647	Genbank	2173834
Sequence 14353	E05957	Genbank	2174144
Sequence 14354	E08663	Genbank	2176776
Sequence 14355	E12274	Genbank	3251108
Sequence 14356	E12457	Genbank	3251290
Sequence 14357	E12651	Genbank	3251483
Sequence 14358	E13330	Genbank	3252135
Sequence 14359	E13405	Genbank	3252210
Sequence 14360	J00139	Genbank	182715
Sequence 14361	J01415	Genbank	1944628
Sequence 14362	J02642		
		Genbank Canbank	182862
Sequence 14363	J02960	Genbank	178203
Sequence 14364	J03248	Genbank	183053
Sequence 14365	J03459	Gen <b>ban</b> k	187172

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Sequence 14366	J03537	Genbank	337513
Sequence 14367	J03592	Genbank	339722
Sequence 14368	J03746	Genbank	183655
Sequence 14369	J03779	Genbank	179833
Sequence 14370	J03934	Genbank	189245
Sequence 14371	J04173	Genbank	551173
Sequence 14372	J04801	Genbank	190371
Sequence 14373	J04991	Genbank	189425
Sequence 14374	J05633	Genbank	186504
Sequence 14375	K00558	Genbank	340020
Sequence 14376	K03432	Genbank	337377
Sequence 14377	L01042	Genbank	184097
Sequence 14378	L01057	Genbank	292462
Sequence 14379	L02320	Genbank	307365
Sequence 14379	L05094	Genbank	388033
Sequence 14381	L05491	Genbank	292831
Sequence 14382	L05628	Genbank	1835658
		Genbank	292442
Sequence 14383	L06498	Genbank	292434
Sequence 14384	L07287	Genbank Genbank	
Sequence 14385	L07633		186512
Sequence 14386	L08048	Genbank	184250
Sequence 14387	L08441	Genbank	179295
Sequence 14388	L09054	Genbank	307344
Sequence 14389	L10284	Genbank	186522
Sequence 14390	L10612	Genbank	402701
Sequence 14391	L11066	Genbank	307322
Sequence 14392	L11370	Genbank	387674
Sequence 14393	L12136	Genbank	181536
Sequence 14394	L12387	Genbank	459835
Sequence 14395	L12711	Genbank	388890
Sequence 14396	L13278	Genbank	292414
Sequence 14397	L14778	Genbank	306476
Sequence 14398	L19185	Genbank	440307
Sequence 14399	L19437	Genbank	4995970
Sequence 14400	L19739	Genbank	431318
Sequence 14401	L19783	Genbank	404725
Sequence 14402	L20046	Genbank	306741
Sequence 14403	L20941	Genbank	507251
Sequence 14404	L22569	Genbank	348706
Sequence 14405	L24444	Genbank Genbank	495860
Sequence 14406	L25931	Genbank Genbank	438638
Sequence 14407	L28809	Genbank Genbank	454151
Sequence 14408	L28997		607027
Sequence 14409	L29766	Genbank	537525 1220360
Sequence 14410	L31610	Genbank	
Sequence 14411	L37127	Genbank Canbank	4164098
Sequence 14412	L38995	Genbank	704415
Sequence 14413	L39320	Genbank	1226044
Sequence 14414	L39326	Genbank	1228094
Sequence 14415	L40396	Genbank	887357
Sequence 14416	L43575	Genbank	899064
Sequence 14417	L43964	Genbank	951202
Sequence 14418	L44140	Genbank	1203968
Sequence 14419	L47162	Genbank	1082035
Sequence 14420	L47647	Genbank	1000861
Sequence 14421	L76159	Gen <b>ban</b> k	1246232

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Sequence 14422	L76416	Genbank	5566605
Sequence 14423	M10036	Genbank	339840
Sequence 14424	M11146	Genbank	182504
Sequence 14425	M11147	Genbank	182513
Sequence 14426	M11353	Genbank	184092
Sequence 14-127	M12623	Genbank	184233
Sequence 14428	M15205	Genbank	339718
Sequence 14429	M15796	Genbank	181271
Sequence 14430	M16541	Genbank	1280204
Sequence 14431	M17886	Genbank	190233
Sequence 14432	M18048	Genbank	337699
Sequence 14433	M19997	Genbank	181968
Sequence 14434	M21142	Genbank	183402
Sequence 14435	M21575	Genbank	1311702
Sequence 14436	M21895	Genbank	189523
Sequence 14437	M22760	Genbank	695359
Sequence 14438	M22918	Genbank	189019
Sequence 14439	M22920	Genbank	189021
	M23410	Genbank	762884
Sequence 14440	M24070	Genbank	181485
Sequence 14441		Genbank	
Sequence 14442	M24194		187701
Sequence 14443	M24486	Genbank	190785
Sequence 14444	M24543	Genbank	341200
Sequence 14445	M24690	Genbank	341218
Sequence 14446	M26481	Genbank	619789
Sequence 14447	M26663	Genbank	618463
Sequence 14448	M29064	Genbank	337452
Sequence 14449	M29960	Genbank	339886
Sequence 14450	M31468	Genbank	190876
Sequence 14451	M31520	Genbank	337504
Sequence 14452	M36341	Genbank	178984
Sequence 14453	M36981	Genbank	189239
Sequence 14454	M37104	Genbank	179274
Sequence 14455	M37583	Genbank	184059
Sequence 14456	M38188	Genbank	189378
Sequence 14457	M55150	Genbank	182392
Sequence 14458	M55409	Genbank	189596
Sequence 14459	M58485	Genbank	180154
Sequence 14460	M60457	Genbank	181249
Sequence 14461	M60484	Genbank	190225
Sequence 14462	M60756	Genbank	184085
Sequence 14463	M60857	Genbank	181334
Sequence 14464	M61764	Genbank	183702
Sequence 14465	M63180	Genbank	339679
Sequence 14466	M73547	Genbank	190161
Sequence 14467	M74775	Genbank	187151
Sequence 14468	M75099	Genbank	337369
Sequence 14469	M76299	Genbank	188778
Sequence 14470	M77232	Genbank	307392
Sequence 14471	M86181	Genbank	190566
Sequence 14472	M87339	Genbank	1498255
Sequence 14473	M88006	Genbank	178522
Sequence 14474	M88468	Genbank	307197
Sequence 14475	M90357	Genbank	179577
Sequence 14476	M93036	Genbank	182904
Sequence 14477	M93651	Genbank	338038

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<u>Table 3-5</u>

Sequence 14478	M94556	Genbank	188855
Sequence 14479	M96982	Genbank	338262
Sequence 14480	S48220	Genbank	257451
Sequence 14481	S54761	Genbank	265221
Sequence 14482	S72871	Genbank	639594
Sequence 14483	S73498	Genbank	688010
Sequence 14484	S73591	Genbank	688296
Sequence 14485	S74678	Genbank	-
Sequence 14486	S81752		241477
		Genbank	1438780
Sequence 14487	S82597	Genbank	1699343
Sequence 14488	S82616	Genbank	1699307
Sequence 14489	U00238	Genbank	404860
Sequence 14490	U02031	Genbank	451329
Sequence 14491	U03886	Genbank	458225
Sequence 14492	U07231	Genbank	517195
Sequence 14493	U07857	Genbank	469048
Sequence 14494	U09466	Genbank	495492
Sequence 14495	U09813	Genbank	1008454
Sequence 14496	U10248	Genbank	984280
Sequence 14497	U10362	Genbank	505651
Sequence 14498	U12404	Genbank	531170
Sequence 14499	U12465	Genbank	562073
Sequence 14500	U12897	Genbank	5 <b>62075</b>
Sequence 14501	U13616	Genbank	608024
Sequence 14502	U14193	Genbank	55 <b>5899</b>
Sequence 14503	U14970	Genbank	550020
Sequence 14504	U14972	Genbank	550024
Sequence 14505	U14973	Genbank	550026
Sequence 14506	U17899	Genbank	717053
Sequence 14507	U18321	Genbank	603763
Sequence 14508	<b>U1867</b> 1	Genbank	1293919
Sequence 14509	U20285	Genbank	64487 <b>8</b>
Sequence 14510	U20362	Genbank	755 <b>485</b>
Sequence 14511	U25182	Genbank	799380
Sequence 14512	U29171	Genbank	881618
Sequence 14513	U29589	Genbank	903978
Sequence 14514	U30826	Genbank	1049079
Sequence 14515	U34683	Genbank	1236349
Sequence 14516	U39817	Genbank	1072121
Sequence 14517	U41060	Genbank	1256000
Sequence 14518	U41195	Genbank	1244694
Sequence 14519	U41448	Genbank	3335023
Sequence 14520	U41668	Genbank	1477481
Sequence 14521	U44798	Genbank	1174216
Sequence 14522	U46751	Genbank	3077821
Sequence 14523	U49245	Genbank	1216503
Sequence 14524	U51559	Genbank	1236709
Sequence 14525	U52111	Genbank	8331754
Sequence 14526	U52852	Genbank	8370402
Sequence 14527	U54993	Genbank	4097314
Sequence 14528	U56833	Genbank	1465750
Sequence 14529	U57846	Genbank	1373418
Sequence 14530	U57847	Genbank	1373416
Sequence 14531	U59911	Genbank	1403712
Sequence 14532	U66243	Genbank	
Sequence 14532	U66589	Genbank	1772645
ocque loc 14000	000003	Genbank	1575566

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<u>Table 3-5</u>

Sequence 14534	U67243	Genbank	2228516
Sequence 14535	U68566	Genbank	1916621
Sequence 14536	U68758	Genbank	4097815
Sequence 14537	U72648	Genbank	3914602
Sequence 14538	U73168	Genbank	1613899
Sequence 14539	U73645	Genbank	1737200
Sequence 14540	U73824	Genbank	1857236
Sequence 14541	U75285	Genbank	2315862
Sequence 14542	U76549	Genbank	1673574
Sequence 14543	U76669	Genbank	6672211
Sequence 14544	U77643	Genbank	2062390
Sequence 14545	U80017	Genbank	1737211
		Genbank	2341006
Sequence 14546	U80460		
Sequence 14547	U85625	Genbank	5091494
Sequence 14548	U87460	Genbank	2076881
Sequence 14549	U90904	Genbank	1913882
Sequence 14550	U90907	Genbank	1913886
Sequence 14551	U91327	Genbank	1871209
Sequence 14552	U91328	Genbank	2088550
Sequence 14553	U94586	Genbank	1946 <b>691</b>
Sequence 14554	U95689	Genbank	<b>2804669</b>
Sequence 14555	U96629	Genbank	2341008
Sequence 14556	V00478	Genbank	28244
Sequence 14557	V00662	Genbank	13003
Sequence 14558	X00351	Genbank	2 <b>82</b> 51
Sequence 14559	X02152	Genbank	3 <b>43</b> 12
Sequence 14560	X03083	Genbank	3 <b>4309</b>
Sequence 14561	X03205	Genbank	36162
Sequence 14562	X05276	Genbank	37201
Sequence 14563	X05332	Genbank	35740
Sequence 14564	X05857	Genbank	31988
Sequence 14565	X06409	Genbank	30204
Sequence 14566	X14608	Genbank	296365
Sequence 14567	X15822	Genbank	30146
Sequence 14568	X15940	Genbank	36129
Sequence 14569	X16135	Genbank	32355
Sequence 14570	X16560	Genb <b>ank</b>	30154
Sequence 14571	X17025	Genbank	488749
Sequence 14572	X17620	Genbank	35067
Sequence 14573	X52839	Genbank	36125
Sequence 14574	X52851	Genba <b>nk</b>	30167
Sequence 14575	X52882	Genbank	311380
Sequence 14576	X52897	Genbank	24484
Sequence 14577	X52943	Genbank	28912
Sequence 14578	X56932	Genbank	23690
Sequence 14579	X56998	Genbank	37564
Sequence 14580	X58288	Genbank	32455
Sequence 14581	X59434	Genbank	432375
Sequence 14582	X60221	Genbank	509290
Sequence 14583	X61755	Genbank	32387
Sequence 14584	X63432	Genbank	28335
Sequence 14585	X63657	Genbank	296185
Sequence 14586	X64123	Genbank	35819
Sequence 14587	X65923	Genbank	31302
	X66975	Genbank	32353
Sequence 14588			3614 <b>9</b>
Sequence 14589	X67247	Genbank	30149

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Sequence 14590	X6769 <b>8</b>	Genbank	37476
Sequence 14591	X69910	Gerbank	297407
Sequence 14592	X7C940	Gerbank	38455
Sequence 14593	X71428	Gerbank	393415
Sequence 14594	X72841	Genbank	297903
Sequence 14595	X74801	Genbank	671526
Sequence 14596	X74979	Genbank	400462
Sequence 14597	X75593	Genbank	452319
Sequence 14598	X75684	Genbank	429092
Sequence 14599	X75861	Genbank	456258
Sequence 14600	X76013	Genbank	531595
Sequence 14601	X78137	Genbank	460770
Sequence 14602	X80200	Genbank	951276
Sequence 14603	X81713	Genbank	558213
Sequence 14604	X81788	Genbank	1045058
Sequence 14605	X81895	Genbank	2370158
Sequence 14606	X81900	Genbank	2274973
Sequence 14607	X84908	Genbank	1502344
Sequence 14608	X85372	Genbank	
Sequence 14609	X87241	Genbank	806563
Sequence 14610	X87344	Genbank	1107686 1054740
Sequence 14611	X87949	Genbank	
Sequence 14612	X91249	Genbank	1143491
Sequence 14613	X93036	Genbank	1160185
Sequence 14614	X94910	Genbank	1085025
Sequence 14615	X95592	Genbank	3413292
Sequence 14616	X96586	Genbank	1185118
Sequence 14617	X97544	Genbank	1556398 1770563
Sequence 14618	×98356	Genbank	
Sequence 14619	X99585	Genbank	1648869
Sequence 14620	Y00052	Genbank	1770518
Sequence 14621	Y00282	Genbank	30308
Sequence 14622	Y00345	Genbank	36048
Sequence 14623	Y00815	Genbank	35569
Sequence 14624	Y08201	Genbank	34266
Sequence 14625	Y09836	Genbank	1552548 1743253
Sequence 14626	Y10345	Genbank	
Sequence 14627	Y15286	Genbank	2292905 2584788
Sequence 14628	Z11692	Genbank	31107
Sequence 14629	Z11793	Genbank	36425
Sequence 14630	Z13009	Genbank	31072
Sequence 14631	Z14244	Genbank	30150
Sequence 14632	Z21507	Genbank	38521
Sequence 14633	Z24724	Genbank	505034
Sequence 14634	Z25749	Genbank	550116
Sequence 14635	Z31695	Genbank	469143
Sequence 14636	Z36832	Genbank	533946
Sequence 14637	Z36849	Genbank	
Sequence 14638	Z46973	Genbank	533963 987947
Sequence 14639	Z47087	Genbank	
Sequence 14640	Z48950	Genbank	860989 761715
Sequence 14641	Z54246	Genbank Genbank	761715
Sequence 14642	Z68164		1000700
Sequence 14643	Z68284	Genbank	1107692
Sequence 14644	Z69711	Genbank Genbank	1130698
Sequence 14645	Z69890		1204109
22400100 14040	703020	Genbank	1213361

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Sequence 14646	Z73900	Genbank	1340069
Sequence 14647	Z82242	Genbank	1841915
Sequence 14648	Z82975	Genbank	1673497
Sequence 14649	Z84474	Genbank	1903188
Sequence 14650	Z84812	Genbank	1845164
		Genbank	4034056
Sequence 14651	Z85986		
Sequence 14652	Z92544	Genbank	1869775
Sequence 14653	Z92845	Genbank	2462406
Sequence 14654	Z93041	Genbank	1905788
Sequence 14655	Z93928	Genbank	2578051
Sequence 14656	Z95113	Genbank	6572188
Sequence 14657	Z95126	Genbank	2342581
Sequence 14658	Z95704	Genbank	2121307
Sequence 14659	Z97053	Genbank	9650676
	Z97054	Genbank	2370077
Sequence 14660		Genbank	2832593
Sequence 14661	Z97056		4455632
Sequence 14662	Z97353	Genbank	_
Sequence 14663	Z97652	Genbank	6456822
Sequence 14664	Z98049	Genbank	2462401
Sequence 14665	Z98749	Genbank	4775611
Sequence 14666	Z98884	Genbank	5304861
Sequence 14667	Z98950	Genbank	3036783
Sequence 14668	Z99128	Genbank	3135974
Sequence 14669	Z99916	Genbank	4158122
Sequence 14670	AA004759	EST	1448327
Sequence 14671	AA007319	EST	1463464
Sequence 14672	AA009421	EST	1470619
	AA009598	EST	1470757
Sequence 14673			1471099
Sequence 14674	AA010071	EST	
Sequence 14675	AA010798	EST	1471844
Sequence 14676	AA011369	EST	1472466
Sequence 14677	AA011539	EST	1472565
Sequence 14678	AA011569	EST	1472595
Sequence 14679	AA013065	EST	1474101
Sequence 14680	AA018720	EST	1481994
Sequence 14681	AA019036	EST	1482627
Sequence 14682	AA019504	EST	1482187
Sequence 14683	AA020851	EST	1484622
Sequence 14684	AA021306	EST	1484995
Sequence 14685	AA024597	EST	1489502
Sequence 14686	AA025057	EST	1489962
	AA025122	EST	1490037
Sequence 14687	AA026215	EST	1492901
Sequence 14688		EST	1492253
Sequence 14689	AA026316		1492674
Sequence 14690	AA026353	EST	
Sequence 14691	AA026643	EST	1492801
Sequence 14692	AA026758	EST	149255 <b>6</b>
Sequence 14693	AA027229	EST	1492149
Sequence 14694	AA029679	EST	1497082
Sequence 14695	AA034414	EST	1506241
Sequence 14696	AA035260	EST	1506772
Sequence 14697	AA035377	EST	1507101
Sequence 14698	AA037143	EST	1512251
Sequence 14699	AA039589	EST	1516014
Sequence 14700	AA039683	EST	1516155
	AA039734	EST	1516236
Sequence 14701	AAU39734	ESI	1010200

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Secuence 14702	AA039876	EST	1516172
Secuence 14703		EST	1516206
Secuence 14704		EST	1516765
Secuence 14705		EST	1522583
Sequence 14706		EST	1521111
Sequence 14707		EST	1521472
Sequence 14708		EST	1523043
Sequence 14709		EST	1523038
Sequence 14710		EST	1524439
Sequence 14711		EST	1524915
Sequence 14712		EST	1524763
Sequence 14713		EST	1524944
Sequence 14714		EST	1544534
Sequence 14715		EST	1544797
Sequence 14716		EST	1544870
Sequence 14717		EST	1545004
Sequence 14718		EST	1545396
Sequence 14719		EST	1548444
Sequence 14720		EST	1549991
Sequence 14721	AA057541	EST	1550246
Sequence 14722		EST	1550199
Sequence 14723		EST	1556632
Sequence 14724	AA063612	EST	1557579
Sequence 14725		EST	1929208
Sequence 14726		EST	1576401
Sequence 14727	AA069422	EST	1576781
Sequence 14728	AA069655	EST	1577014
Sequence 14729	AA070277	EST	1577637
Sequence 14730	AA070855	EST	1578487
Sequence 14731	AA071269	EST	1578757
Sequence 14732	AA071402	EST	1578773
Sequence 14733	AA071472	EST	1578903
Sequence 14734	AA074221	EST	1614090
Sequence 14735	AA074611	EST	1614677
Sequence 14736	AA074614	EST	1614492
Sequence 14737	AA074664	EST	1614608
Sequence 14738	AA074831	EST	1614875
Sequence 14739	AA075200	EST	1615070
Sequence 14740 Sequence 14741	AA075981	EST	1615850
Sequence 14741	AA079050 AA081478	EST EST	1617942
Sequence 14743	AA082545	EST	162 <b>3536</b> 162 <b>4602</b>
Sequence 14743	AA083712	EST	1625772
Sequence 14745	AA084678	EST	1626735
Sequence 14746	AA084868	EST	1626988
Sequence 14747	AA088189	EST	1633764
Sequence 14748	AA090105	EST	1636589
Sequence 14749	AA092310	EST	1637531
Sequence 14750	AA092324	EST	1637545
Sequence 14751	AA098978	EST	1644862
Sequence 14752	AA099015	EST	1644895
Sequence 14753	AA099080	EST	1645522
Sequence 14754	AA099327	EST	1646033
Sequence 14755	AA100639	EST	1646940
Sequence 14756	AA101116	EST	1647692
Sequence 14757	AA101598	EST	1648552
			_

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Sequence 14758	AA101663	EST	1648744
Sequence 14759	AA102442	EST	1647356
Sequence 14760	AA102560	EST	1647752
Sequence 14761	AA114042	EST	1667919
Sequence 14762	AA115474	EST	1670145
Sequence 14763	AA126478	EST	1686071
Sequence 14764	AA126762	EST	1686262
Sequence 14765	AA126841	EST	1686323
Sequence 14765	AA127483	EST	1686755
Sequence 14767	AA127550	EST	1686822
Sequence 14768	AA127886	EST	1687193
Sequence 14769	AA129205	EST	1688990
Sequence 14770	AA129465	EST	1689336
Sequence 14770	AA129935	EST	1689511
	AA131336	EST	1692834
Sequence 14772	AA132089	EST	1693579
Sequence 14773	AA132394	EST	1693885
Sequence 14774	AA132989	EST	1694505
Sequence 14775	AA133031	EST	1694539
Sequence 14776	AA133736	EST	1690768
Sequence 14777		EST	1690727
Sequence 14778	AA133759 AA134263	EST	1691637
Sequence 14779		EST	1695597
Sequence 14780	AA134609	EST	- 1696141
Sequence 14781	AA135030	EST	1696227
Sequence 14782	AA135126 AA135254	EST	1696373
Sequence 14783	AA136337	EST	1697545
Sequence 14784 Sequence 14785	AA136723	EST	1697933
Sequence 14786	AA136761	EST	1698008
Sequence 14787	AA137016	EST	1698225
Sequence 14788	AA142942	EST	1712320
Sequence 14789	AA143324	EST	1712695
Sequence 14790	AA143422	EST	1712852
Sequence 14791	AA147059	EST	1716433
Sequence 14792	AA147948	EST	1717455
Sequence 14793	AA148889	EST	1718998
Sequence 14794	AA149404	EST	1719920
Sequence 14795	AA149518	EST	1720336
Sequence 14796	AA149579	EST	1720380
Sequence 14797	AA149801	EST	1720899
Sequence 14798	AA149826	EST	1720961
Sequence 14799	AA150361	EST	1721873
Sequence 14800	AA150481	EST	1721995
Sequence 14801	AA151140	EST	1719383
Sequence 14802	AA151423	EST	1719820
Sequence 14803	AA151795	EST	1720490
Sequence 14804	AA151809	EST	1720504
Sequence 14805	AA151816	EST	1720511
Sequence 14806	AA152179	EST	1721231
Sequence 14807	AA155999	EST	1727666
Sequence 14808	AA156226	EST	1727844
Sequence 14809	AA158153	EST	1732990
Sequence 14810	AA158669	EST	1733480
Sequence 14811	AA158704	EST	1733533
Sequence 14812	AA159176	EST	1733969
Sequence 14813	AA159347	EST	1734149

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Sequence 14814	AA160666	EST	1736043
Sequence 14815	AA161467	EST	1735906
Sequence 14816	AA165437	EST	1741453
Sequence 14817	AA166691	EST	1745155
Sequence 14818	AA166843	EST	1745024
Sequence 14819	AA165946	EST	1745322
Sequence 14820	AA167011	EST	1745386
Sequence 14821	AA167501	EST	1745886
Sequence 14822	AA169 <b>786</b>	EST	1748121
Sequence 14823	AA171706	EST	1750791
Sequence 14824	AA173421	EST	1753550
Sequence 14825	AA173705	EST	1753883
Sequence 14826	AA173936	EST	17540 <b>68</b>
Sequence 14827	AA176244	EST	1757375
Sequence 14828	AA176455	EST	1757713
Sequence 14829	AA178955	EST	17603 <b>08</b>
Sequence 14830	AA179220	EST	1760572
Sequence 14831	AA181 <b>891</b>	EST	17653 <b>68</b>
Sequence 14832	AA182751	EST	17665 <b>81</b>
Sequence 14833	AA182940	EST	1 <b>766108</b>
Sequence 14834	AA186328	EST	1774446
Sequence 14835	AA186811	EST	17751 <b>03</b>
Sequence 14836	AA190408	EST	1779239
Sequence 14837	AA192099	EST	1781923
Sequence 14838	AA192347	EST	1781585
Sequence 14839	AA194517	EST	1784213
Sequence 14840	AA195093	EST	1784783
Sequence 14841	AA203477	EST	1799204
Sequence 14842	AA204856	EST	1802881
Sequence 14843	AA2C5970	EST	1801341
Sequence 14844	AA206446	EST	1801826
Sequence 14845	AA2C6761	EST	1802148
Sequence 14846 Sequence 14847	AA210909	EST	1809555
Sequence 14848	AA214548	EST	1813173
Sequence 14849	AA215640 AA216047	EST	1815385
Sequence 14850	AA216047 AA216061	EST	1816018
Sequence 14851	AA219137	EST	1816048
Sequence 14852	AA220941	EST EST	1833202
Sequence 14853	AA223990	EST	1839687
Sequence 14854	AA224985	EST	18445 <b>1</b> 5 1846276
Sequence 14855	AA225438	EST	1846765
Sequence 14856	AA225485	EST	1846812
Sequence 14857	AA225653	EST	1846970
Sequence 14858	AA225814	EST	1847200
Sequence 14859	AA226768	EST	1848466
Sequence 14860	AA227285	EST	1848839
Sequence 14861	AA229713	EST	1851876
Sequence 14862	AA232183	EST	1855538
Sequence 14863	AA235835	EST	1860273
Sequence 14864	AA243394	EST	1874187
Sequence 14865	AA243646	EST	1874456
Sequence 14866	AA243774	EST	1874594
Sequence 14867	AA244401	EST	1875185
Sequence 14868	AA247569	EST	1879338
Sequence 14869	AA251128	EST	1886090

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Sequence 14870	AA252998	EST	1882921
Sequence 14871	AA257006	EST	1891135
Sequence 14872	AA258030	EST	1894480
Sequence 14873	AA258271	EST	1893453
Sequence 14874	AA258394	EST	1893536
Sequence 14875	AA261894	EST	1897728
Sequence 14876	AA261960	EST	1898149
Sequence 14877	AA262181	EST	1898452
Sequence 14878	AA262425	EST	1897921
Sequence 14879	AA262871	EST	1898592
Sequence 14880	AA263047	EST	1898845
	AA278243	EST	1920183
Sequence 14881		EST	1920103
Sequence 14882	AA278787		
Sequence 14883	AA279139	EST	1920605
Sequence 14884	AA279421	EST	1920886
Sequence 14885	AA280206	EST	1921881
Sequence 14886	AA280658	EST	1923453
Sequence 14887	AA280848	EST	1923546
Sequence 14888	AA282253	EST	1925187
Sequence 14889	AA285040	EST	1927721
Sequence 14890	AA285126	EST	1928107
Sequence 14891	AA292069	EST	1940055
Sequence 14892	AA292296	EST	1940276
Sequence 14893	AA294863	EST	19472 <b>08</b>
Sequence 14894	AA295158	EST	1947513
Sequence 14895	AA296277	EST	1948652
Sequence 14896	AA296690	EST	1949182
Sequence 14897	AA297999	EST	1950582
Sequence 14898	AA300796	EST	1953128
Sequence 14899	AA303856	EST	1956421
Sequence 14900	AA304244	EST	1956597
Sequence 14901	AA304272	EST	1956615
Sequence 14902	AA305667	EST	1958242
Sequence 14903	AA306121	EST	195 <b>8449</b>
Sequence 14904	AA306470	EST	1958798
Sequence 14905	AA307154	EST	1959482
Sequence 14906	AA307818	EST	1960145
Sequence 14907	AA308297	EST	1960626
Sequence 14908	AA308596	EST	1961163
Sequence 14909	AA311749	EST	1964076
Sequence 14910	AA313317	EST	1965646
Sequence 14911	AA313802	EST	1966131
Sequence 14912	AA314008	EST	1966337
Sequence 14913	AA314493	EST	1966822
Sequence 14914	AA314869	EST	1967218
Sequence 14915	AA315072	EST	1967411
Sequence 14916	AA315192	EST	1967541
Sequence 14917	AA316110	EST	1968439
Sequence 14918	AA316272	EST	1968601
Sequence 14919	AA316868	EST	1969196
Sequence 14919	AA317537	EST	1969886
Sequence 14921	AA318361	EST	1970709
Sequence 14921	AA319855	EST	1972181
Sequence 14923	AA325081	EST	1977579
Sequence 14923	AA328973	EST	1981216
			1982040
Sequence 14925	AA329553	EST	1302040

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Sequence 1	4926	AA334936	EST	1987254
Sequence 1		AA339220	EST	1991456
Sequence 1		AA341254	EST	1993491
Sequence 1		AA342547	EST	1994804
Sequence 1		AA343888	EST	1996146
Sequence 1		AA343129	EST	2000440
Sequence 1		AA355326		
			EST	2007708
Sequence 1		AA359995	EST	2012346
Sequence 1		AA362311	EST	2014781
Sequence 1		AA362547	EST	2014916
Sequence 1		AA363030	EST	2015589
Sequence 1		AA363971	EST	2016289
Sequence 1		AA364506	EST	2016845
Sequence 1		AA371595	EST	2024137
Sequence 1		AA371677	EST	2024220
Sequence 1		AA371861	EST	2024387
Sequence 1		AA376859	EST	2029176
Sequence 1		AA377534	EST	2029863
Sequence 1	4944	AA378981	EST	2031371
Sequence 1	4945	AA379089	EST	2031419
Sequence 1		AA382161	EST	2034479
Sequence 1	4947	AA382953	EST	2035271
Sequence 1-	4948	AA393055	EST	2046151
Sequence 1-	4949	AA393236	EST	2046205
Sequence 14	4950	AA397442	EST	2050483
Sequence 1	4951	AA398254	EST	2051363
Sequence 14	4952	AA400319	EST	2054395
Sequence 14	4953	AA400551	EST	2054422
Sequence 1	4954	AA400632	EST	2054503
Sequence 14		AA401792	EST	2057276
Sequence 14	4956	AA402613	EST	2056402
Sequence 14	4957	AA402800	EST	2056553
Sequence 14	4958	AA404535	EST	2059268
Sequence 14		AA404961	EST	2063336
Sequence 14		AA405814	EST	2063797
Sequence 14		AA406106	EST	2064105
Sequence 14		AA406543	EST	2064657
Sequence 14		AA410626	EST	2069851
Sequence 14		AA411013	EST	2070137
Sequence 14		AA411736	EST	2069397
Sequence 14		AA412286	EST	2070857
Sequence 14	4967	AA412695	EST	2071301
Sequence 14		AA417129	EST	2077272
Sequence 14		AA417185	EST	2077292
Sequence 14		AA418317	EST	2080146
Sequence 14		AA418533	EST	2080314
Sequence 14		AA418734	EST	2080535
Sequence 14		AA419146	EST	2078944
Sequence 14		AA420564	EST	2094461
Sequence 14		AA420675	EST	2094553
Sequence 14	-	AA420705	EST	2094611
Sequence 14		AA420873	EST	2094306
Sequence 14		AA421675	EST	2100533
Sequence 14		AA422114	EST	2100938
Sequence 14		AA424037	EST	2103034
Sequence 14		AA425108		
padingues 14	+301	MM423100	EST	2107178

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Sequence 14982	AA425723	EST	2106443
Sequence 14983	AA426125	EST	2106597
Sequence 14984	AA426618	EST	2107298
Sequence 14985	AA427438	EST	2111308
Sequence 14986	AA427994	EST	2112068
Sequence 14987	AA427999	EST	2112073
Sequence 14988	AA428399	EST	2110228
Sequence 14989	AA431300	EST	2115008
Sequence 14990	AA431320	· EST	2115028
Sequence 14991	AA431322	EST	2115030
Sequence 14992	AA431581	EST	2115289
Sequence 14993	AA431812	EST	2115520
Sequence 14994	AA432282	EST	2114670
Sequence 14995	AA434140	EST	2139054
Sequence 14996	AA436116	EST	2141030
Sequence 14997	AA436991	EST	2141905
Sequence 14998	AA437166	EST	2142080
Sequence 14999	AA437224	EST	2142138
Sequence 15000	AA442133	EST	2154011
Sequence 15001	AA443297	EST	2155 <b>972</b>
Sequence 15002	AA443762	EST	2156437
Sequence 15003	AA446569	EST	2159234
Sequence 15004	AA447974	EST	2161644
Sequence 15005	AA448292	EST	2161962
Sequence 15006	AA451954	EST	2165623
Sequence 15007	AA453183	EST	2166852
Sequence 15008	AA456223	EST	2179433
Sequence 15009	AA456863	EST	2179583
Sequence 15010	AA458476	EST	2183383
Sequence 15011	AA459991	EST	2184875
Sequence 15012	AA461239	EST	2186359
Sequence 15013	AA461254	EST	2186374
Sequence 15014	AA463447	EST	2188331
Sequence 15015	AA463899	EST	2188783
Sequence 15016	AA464099	EST	2188983
Sequence 15017	AA464353	EST	2189237
Sequence 15018	AA465191	EST	2191358
Sequence 15019	AA465537	EST	2191704
Sequence 15020	AA468049	EST	2194583
Sequence 15021	AA468360	EST	2194894
Sequence 15022	AA470144	EST	2197453
Sequence 15023	AA477119	EST	2205803
Sequence 15024	AA478414	EST	2207048
Sequence 15025	AA478637	EST	2207271
Sequence 15026	AA479488	EST	2208044
Sequence 15027	AA479863	EST	2205749
Sequence 15028	AA479892	EST	2204374
Sequence 15029	AA481045	EST	2210597
Sequence 15030	AA483162	EST	2211975
Sequence 15031	AA483450	EST	2212263
Sequence 15032	AA484155	EST	2212968
Sequence 15033	AA487382	EST	2217546
Sequence 15034	AA487628	EST	2217792
Sequence 15035	AA487671	EST	2217835
Sequence 15036	AA488200	EST	2215631
Sequence 15037	AA488782	EST	2218384

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Sequence 150	038 AA	492042	EST	2221604
Sequence 150		492143	EST	2221705
Sequence 150		492267	EST	2221829
Sequence 150	)41 A/	492423	EST	2221985
Sequence 150		493364	EST	2223205
Sequence 150		493382	EST	2223223
Sequence 150		493522	EST	2223363
Sequence 150		493624	EST	2223465
Sequence 150		493807	EST	2223648
Sequence 150		496002	EST	2229323
Sequence 150		\502198	EST	2237165
Sequence 150	-	NS02130 NS03141	EST	2238108
Sequence 150		1503141 1503404	EST	2238371
Sequence 150		\503941	EST	2238908
		1504095	EST	2240255
Sequence 150		4504531	EST	2240233
Sequence 150				
Sequence 150		\504703	EST	2240863
Sequence 150	755 AF	505440	EST	2241577
Sequence 150		506953	EST	2243392
Sequence 150		508597	EST	2246100
Sequence 150		513357	ES⊤	2251769
Sequence 150		513367	EST	2251779
Sequence 150		514331	EST	2253839
Sequence 150		514474	EST	2254074
Sequence 150		514538	EST	2254138
Sequence 150		515736	EST	2255336
Sequence 150		520996	EST	2261539
Sequence 150		521142	EST	2261685
Sequence 150		521353	EST	2261896
Sequence 150		522584	EST	2263296
Sequence 150	68 AA	523005	EST	2388772
Sequence 150		524825	EST	2265753
Sequence 150		525374	EST	2266302
Sequence 150		525960	EST	2 <b>268029</b>
Sequence 150		526186	EST	2268255
Sequence 150	-	526243	EST	2268312
Sequence 150		526569	EST	2268638
Sequence 150		527150	EST	2269219
Sequence 150		527943	EST	2270012
Sequence 150		528066	EST	2270135
Sequence 150		531276	EST	2273982
Sequence 150		531295	EST	2274001
Sequence 150		532807	EST	2278383
Sequence 150		533483	EST	2277579
Sequence 150		534137	EST	2278153
Sequence 150		534 <b>340</b>	EST	2278593
Sequence 150		535571	EST	2279824
Sequence 150		53 <b>5832</b>	EST	2280085
Sequence 150		541343	EST	2287777
Sequence 150		541537	EST	2287971
Sequence 150		541666	EST	2288100
Sequence 150		54 <b>8238</b>	EST	2318520
Sequence 150	90 AA	551141	EST	2321393
Sequence 150	91 AA	551358	EST	2321610
Sequence 150	92 AA	551385	EST	2321637
Sequence 150	93 AA	551581	EST	2321833

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<u>Table 3-5</u>

Sequence 15094	AA551709	EST	2321961
Sequence 15095	AA551728	EST	2321980
Sequence 15095	AA551770	EST	2322022
	AA552169	EST	2322421
Sequence 15097		EST	2322542
Sequence 15098	AA552290	EST	2324564
Sequence 15099	AA554025		2325257
Sequence 15100	AA554718	EST	
Sequence 15101	AA554874	EST	2325413
Sequence 15102	AA555194	EST	2325733
Sequence 15103	AA555326	EST	2325865
Sequence 15104	AA557229	EST	2327706
Sequence 15105	AA557760	EST	2328237
Sequence 15106	AA557768	EST	2328245
Sequence 15107	AA563778	EST	2335417
Sequence 15108	AA566072	EST	2337711
Sequence 15109	AA568232	EST	2341286
Sequence 15110	AA570116	EST	2344096
Sequence 15110	AA570171	EST	2344151
Sequence 15111	AA570649	EST	2344629
Sequence 15112		EST	2344685
Sequence 15113	AA570705	EST	2347843
Sequence 15114	AA573315	EST	2347882
Sequence 15115	AA573354		2348087
Sequence 15116	AA573559	EST	2350454
Sequence 15117	AA575939	EST	2354006
Sequence 15118	AA576506	EST	
Sequence 15119	AA578 <b>782</b>	EST	2356966
Sequence 15120	AA5793 <b>7</b> 5	EST	2357559
Sequence 15121	AA579617	EST	2357801
Sequence 15122	AA579690	EST	2357874
Sequence 15123	AA579834	EST	2355161
Sequence 15124	AA580489	EST	2355816
Sequence 15125	AA582370	EST	2359730
Sequence 15126	AA586421	EST	2398416
Sequence 15127	AA588541	EST	2401716
Sequence 15128	AA593040	EST	2408 <b>802</b>
Sequence 15129	AA595014	EST	2410364
Sequence 15130	AA595339	EST	2410689
Sequence 15131	AA599268	EST	2432893
Sequence 15132	AA599661	EST	2433 <b>286</b>
Sequence 15133	AA600340	EST	24339 <b>65</b>
Sequence 15134	AA600781	EST	2434406
Sequence 15135	AA602357	EST	2436335
Sequence 15136	AA604839	EST	2445 <b>703</b>
Sequence 15137	AA604894	EST	2 <b>445758</b>
Sequence 15138	AA608917	EST	2457345
Sequence 15139	AA608990	EST	2457418
Sequence 15140	AA609219	EST	2457647
Sequence 15141	AA613550	EST	24 <b>64588</b>
Cogneroe 15142	AA613637	EST	2464675
Sequence 15142	AA618493	EST	2505698
Sequence 15143	AA620307	EST	2524246
Sequence 15144	_	EST	2524404
Sequence 15145	AA620465	EST	2525135
Sequence 15146	AA621196	EST	2526682
Sequence 15147	AA622806	EST	2537540
Sequence 15148	AA625155		2537928
Sequence 15149	AA625541	EST	2001 920

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<u>Table 3-5</u>

Sequence 15150	AA625377	EST	2538764
Sequence 15151	AA626635	EST	2539022
Secuence 15152	AA626658	EST	2539045
Secuence 15153	AA628413	EST	
Secuence 15154	AA629923		2540800
Secuence 15155		EST	2552534
_	AA629979	EST	2552590
Sequence 15156	AA630335	EST	2552946
Sequence 15157	AA631646	EST	2554257
Sequence 15158	AA631803	EST	2554414
Sequence 15159	AA631901	EST	2554512
Sequence 15160	AA632126	EST	2555540
Sequence 15161	AA632225	EST	2555639
Sequence 15162	AA632270	EST	2555684
Sequence 15163	AA632610	EST	2556024
Sequence 15164	AA633438	EST	2555298
Sequence 15165	AA633491	EST	2555351
Sequence 15166	AA634087	EST	2557301
Sequence 15167	AA634119	EST	2557333
Sequence 15168	AA635616	EST	2559458
Sequence 15169	AA639362	EST	
Sequence 15170	AA640626	EST	2563141 2565876
Sequence 15171	AA641037		
Sequence 15172	AA641660	EST	2566287
Sequence 15173	AA644009	EST	2566878
Sequence 15174	AA644184	EST	2569227
		EST	2569402
Sequence 15175 Sequence 15176	AA648402 AA649327	EST	2574831
Sequence 15177	AA649926	EST	2575756
Sequence 15177	AA649975	EST	2577254
Sequence 15178		EST	2577303
Sequence 15179	AA651721 AA652395	EST	2583373
Sequence 15181	AA659388	EST	2584047
Sequence 15182	AA659693	EST	2595542
		EST	2595847
Sequence 15183	AA661905	EST	2615996
Sequence 15184	AA662530	EST	2616621
Sequence 15185	AA663778	EST	2617769
Sequence 15186	AA665413	EST	2620026
Sequence 15187	AA669980	EST	2631479
Sequence 15188	AA676967	EST	2657489
Sequence 15189	AA677766	EST	2658288
Sequence 15190	AA679533	EST	2660055
Sequence 15191	AA683488	EST	2670086
Sequence 15192	AA687192	EST	2674397
Sequence 15193	AA687631	EST	2674537
Sequence 15194	AA687667	EST	2674573
Sequence 15195	AA689603	EST	2690530
Sequence 15196	AA699712	EST	2702675
Sequence 15197	AA699807	EST	2702770
Sequence 15198	AA702016	EST	2705129
Sequence 15199	AA702170	EST	2705283
Sequence 15200	AA705025	EST	2714943
Sequence 15201	AA707310	EST	2717228
Sequence 15202	AA708327	EST	2718245
Sequence 15203	AA714016	EST	2726290
Sequence 15204	AA714225	EST	2726499
Seauence 15205	AA716100	EST	2728374

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		FCT	2728588
Sequence 15206	AA716314	EST EST	2736815
Sequence 15207	AA720680	EST	2736925
Sequence 15208	AA720790	EST	2737763
Sequence 15209	AA721701	EST	2740971
Sequence 15210	AA723194	EST	2743179
Sequence 15211	AA725472	EST	2743528
Sequence 15212	AA725821	EST	2750251
Sequence 15213	AA728892	EST	2750320
Sequence 15214	AA728961	EST	2750740
Sequence 15215	AA729381	EST	2752332
Sequence 15216	AA731128 AA731392	EST	2753548
Sequence 15217	AA731610	EST	2753766
Sequence 15218	AA733210	EST	2775319
Sequence 15219	AA736880	EST	2768114
Sequence 15220	AA742380	EST	2784380
Sequence 15221	AA742661	EST	2782167
Sequence 15222	AA744728	EST	2783492
Sequence 15223	AA744747	EST	2783511
Sequence 15224	AA744963	EST	2783727
Sequence 15225	AA745648	EST	2785634
Sequence 15226	AA746967	EST	2786925
Sequence 15227 Sequence 15228	AA747290	EST	2787248
Sequence 15229	AA747733	EST	2787691
Sequence 15230	AA749010	EST	2788968
Sequence 15231	AA749335	EST	2789293
Sequence 15232	AA764788	EST	2816026
Sequence 15233	AA765778	EST	2817016
Sequence 15234	AA768205	EST	2819220
Sequence 15235	AA769138	EST	2820376
Sequence 15236	AA769353	EST	2820591 2820776
Sequence 15237	AA769538	EST	2820776 2821224
Sequence 15238	AA769986	EST	2821454
Sequence 15239	AA770216	EST	2821851
Sequence 15240	AA770613	EST	2823483
Sequence 15241	AA771700	EST	2825989
Sequence 15242	AA774100	EST	2834749
Sequence 15243	AA775415	EST EST	2834958
Sequence 15244	AA775624	EST	2837687
Sequence 15245	AA778356	EST	2837991
Sequence 15246	AA778660	EST	2838256
Sequence 15247	AA778925 AA779599	EST	2838930
Sequence 15248	AA780299	EST	2839630
Sequence 15249	AA782060	EST	2841391
Sequence 15250	AA782690	EST	2842021
Sequence 15251	AA805264	EST	2874014
Sequence 15252	AA806729	EST	2875479
Sequence 15253	AA807199	EST	2876775
Sequence 15254	AA807824	EST	2877230
Sequence 15255 Sequence 15256	AA809127	EST	2878533
Sequence 15257	AA810277	EST	2879636
Sequence 15258	AA812398	EST	2882009
Sequence 15259	AA814392	EST	2883988
Sequence 15260		EST	2884324
Sequence 15261	AA814888	EST	2884484
36436100 10501			

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# <u>Table 3-5</u>

Sequence 1526	2 AA815 <b>42</b> 2	EST	2885018
Sequence 1526		EST	2896535
Sequence 1526		EST	2899289
Sequence 1526		EST	2901317
Sequence 1526		EST	2900168
Sequence 1526		EST	2900179
Sequence 1526		EST	2900365
Sequence 15269		EST	2900366
Sequence 1527		EST	
Sequence 1527		EST	2901720
Sequence 1527		EST	2905025
			2905047
Sequence 15270		EST	2898667
Sequence 15274		EST	2908787
Sequence 15275		EST	2910067
Sequence 15276		EST	2910188
Sequence 15277		EST	2912327
Sequence 15278		EST	2913334
Sequence 15279		EST	2929715
Sequence 15280		EST	2930041
Sequence 15281		EST	2931407
Sequence 15282		EST	2932875
Sequence 15283		EST	2932896
Sequence 15284		EST	2932955
Sequence 15285		EST	2941475
Sequence 15286		EST	2946464
Sequence 15287		EST	2953697
Sequence 15288		EST	2955709
Sequence 15289		EST	<b>295</b> 5711
Sequence 15290		EST	2955763
Sequence 15291		EST	2958810
Sequence 15292		EST	2958886
Sequence 15293		EST	2957607
Sequence 15294		EST	2957828
Sequence 15295		EST	2958147
Sequence 15296		EST	2958432
Sequence 15297	' AA868706	EST	2964151
Sequence 15298	AA873173	EST	2969295
Sequence 15299	AA873175	EST	2969297
Sequence 15300	AA877181	EST	2986258
Sequence 15301		EST	29 <b>8</b> 6727
Sequence 15302		EST	2992812
Sequence 15303		EST	2994370
Sequence 15304		EST	2994374
Sequence 15305		EST	3001141
Sequence 15306	AA887219	EST	3002327
Sequence 15307		EST	3017224
Sequence 15308	AA890400	EST	3017279
Sequence 15309	AA890449	EST	3017328
Sequence 15310		EST	3017355
Sequence 15311	AA902 <b>974</b>	EST	3038097
Sequence 15312		EST	3040485
Sequence 15313		EST	3041473
Sequence 15314	AA908188	EST	3047593
Sequence 15315		EST	3052431
Sequence 15316		EST	3058500
Sequence 15317		EST	3075842
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<u>Table 3-5</u>

Sequence 15374	AI073499	EST	34 <b>00143</b>
Sequence 15375	AI078597	EST	341 <b>3005</b>
Sequence 15376	Al081106	EST	341 <b>7898</b>
Sequence 15377	Al081419	EST	3418211
Sequence 15378	Al081478	EST	3418270
		EST	3418328
Sequence 15379	AI081536	EST	3423030
Sequence 15380	A1084607		
Sequence 15381	AI086937	EST	3425 <b>360</b>
Sequence 15382	A1087840	EST	3426873
Sequence 15383	AI089289	EST	3428348
Sequence 15384	AI089629	EST	342 <b>8688</b>
Sequence 15385	AI091340	EST	3430399
Sequence 15386	AI091999	EST	3427197
Sequence 15387	A1092232	EST	3431 <b>226</b>
Sequence 15388	Al092931	EST	3431 <b>907</b>
Sequence 15389	AI093444	EST	3432 <b>420</b>
Sequence 15390	Al094174	EST	3433150
Sequence 15391	A1095534	EST	3434510
Sequence 15392	A1095786	EST	3434762
Sequence 15393	AI096685	EST	3446179
Sequence 15394	AI114501	EST	6359846
Sequence 15395	Al114683	EST	6360 <b>028</b>
Sequence 15396	Al124954	EST	3593 <b>468</b>
Sequence 15397	Al125449	EST	3593 <b>963</b>
Sequence 15398	Al125479	EST	3593 <b>993</b>
Sequence 15399	Al125915	EST	3594 <b>429</b>
Sequence 15400	Al127049	EST	35955 <b>63</b>
Sequence 15401	Al127824	EST	3596 <b>338</b>
Sequence 15402	Al129324	EST	3597838
Sequence 15403	Al129703	EST	3598217
Sequence 15404	Al129827	EST	3598341
Sequence 15405	Al130744	EST	3600760
Sequence 15406	Al130885	EST	3600901
Sequence 15407	Al132921	EST	6360237
Sequence 15408	Al133690	EST	6361006
Sequence 15409	Al138441	EST	3644413
Sequence 15410	Al139393	EST	3645 <b>365</b>
Sequence 15410	Al141041	EST	36484 <b>98</b>
Sequence 15411	Al142953	EST	3659312
Sequence 15412	AI142933 AI143184	EST	3664993
Sequence 15413	Al143226	EST	3665035
Sequence 15414	Al143638	EST	3665447
Sequence 15415	A1143636 A1143894	EST	3665 <b>703</b>
Sequence 15416	AI143694 AI144216	EST	3666025
Sequence 15417		EST	3666194
Sequence 15418	Al144385 Al146512	EST	36741 <b>9</b> 4
Sequence 15419		EST	3679313
Sequence 15420	Al150844		3679347
Sequence 15421	AI150878	EST	369 <b>3252</b>
Sequence 15422	AI159872	EST	3693379
Sequence 15423	Al159999	EST	
Sequence 15424	Al160331	EST	369 <b>3711</b>
Sequence 15425	Al160430	EST	3693810
Sequence 15426	Al161162	EST	369 <b>4467</b>
Sequence 15427	Al167145	EST	3700315
Sequence 15428	Al167693	EST	3700863
Sequence 15429	Al168090	EST	3701 <b>260</b>

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<u>Table 3-5</u>

Sequence 15430	AI174394	EST	3721247
Sequence 15431	AI174722	EST	6361100
Sequence 15432	Al174834	EST	6361229
Sequence 15433	AI188128	EST	3739337
Sequence 15434	Al188187	EST	3739396
Sequence 15435	AI188575	EST	3739784
Sequence 15436	Al188768	EST	3739977
Sequence 15437	AI189569	EST	3740778
Sequence 15438	Al191342	EST	3742551
Sequence 15439	Al191933	EST	
Sequence 15440	Al199229	EST	3743142
Sequence 15441	Al201859		3751835
Sequence 15442	Al201659 Al203123	EST	3754465
		EST	3755729
Sequence 15443	AI207440	EST	6361448
Sequence 15444	Al207448	EST	6361456
Sequence 15445	AI208443	EST	3770385
Sequence 15446	Al218943	EST	3801146
Sequence 15447	AI222237	EST	3804440
Sequence 15448	AI222424	EST	3804627
Sequence 15449	AI222782	EST	3804985
Sequence 15450	Al222806	EST	3805009
Sequence 15451	Al240909	EST	3836306
Sequence 15452	Al241212	EST	3836609
Sequence 15453	Al241763	EST	3837160
Sequence 15454	Al244187	EŞT	3839584
Sequence 15455	Al245887	EST	3841284
Sequence 15456	Al246248	EST	3841645
Sequence 15457	AI248623	EST	3844020
Sequence 15458	Al249146	EST	3844543
Sequence 15459	Al249680	EST	3846209
Sequence 15460	Al253055	EST	3849584
Sequence 15461	Al253330	EST	3850451
Sequence 15462	Al253414	EST	3850369
Sequence 15463	Al253609	EST	3858537
Sequence 15464	Al254360	EST	3861885
Sequence 15465	Al261577	EST	3869780
Sequence 15466	Al263373	EST	3871576
Sequence 15467	Al264005	EST	3872208
Sequence 15468	Al264289	EST	3872492
Sequence 15469	Al266408	EST	3884566
Sequence 15470	Al266746	EST	3884904
Sequence 15471	Al267341	EST	3886508
Sequence 15472	Al267471	EST	3886638
Sequence 15473	Al267960	EST	3887127
Sequence 15474	Al268542	EST	3887709
Sequence 15475	Al269537	EST	3888704
Sequence 15476	Al271234	EST	3890401
Sequence 15477	Al271608	EST	3890775
Sequence 15478	Al274628	EST	3896896
Sequence 15479	Al276950	EST	3899218
Sequence 15480	Al278211	EST	3900479
Sequence 15481	Al278317	EST	3900585
Sequence 15482	Al278411	EST	3900679
Sequence 15483	Al278998	EST	3917232
Sequence 15484	Al279500	EST	3917734
Sequence 15485	Al279903	EST	3918137
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Sequence 15486	Al281560	EST	3919793
Sequence 15487	Al282246	EST	3920479
Sequence 15488	Al283100	EST	3921333
Sequence 15489	Al284545	EST	3922778
Sequence 15490	Al284565	EST	3922798
Sequence 15491	Al289108	EST	3932372
Sequence 15492	Al289752	EST	3931593
Sequence 15493	Al290104	EST	3931770
Convence 15495	Al291500	EST	3934274
Sequence 15494	Al291613	EST	3934387
Sequence 15495		EST	3959038
Sequence 15496	Al299384		
Sequence 15497	Al299392	EST	3959046
Sequence 15498	Al300346	EST	3959692
Sequence 15499	Al300555	EST	3959901
Sequence 15500	Al301294	EST	3960640
Sequence 15501	Al301904	EST	3961250
Sequence 15502	Al305197	EST	3988838
Sequence 15503	Al305516	EST	3990407
Sequence 15504	Al307736	EST	4001954
Sequence 15505	Al311614	EST	4006485
Sequence 15506	Al313335	EST	4018940
Sequence 15507	Al332968	EST	4069527
Sequence 15508	Al332994	EST	4069553
Sequence 15509	Al334291	EST	4070850
Sequence 15510	Al335537	EST	4072464
Sequence 15511	Al339426	EST	4076353
Sequence 15512	AI339635	EST	4076562
Sequence 15513	Al339952	EST	4076879
Sequence 15514	Al341076	EST	4078003
Sequence 15515	Al342005	EST	4078932
Sequence 15516	Al342217	EST	4079144
Sequence 15517	Al342486	EST	4079413
Sequence 15518	Al344424	EST	4081630
Sequence 15519	Al345709	EST	4082915
Sequence 15520	Al347461	EST	4084667
Sequence 15521	Al348001	EST	4085207
Sequence 15522	Al348771	EST	4085989
Sequence 15523	Al349772	EST	4086978
Sequence 15524	Al352497	EST	4089703
Cocuence 15525	Al354464	EST	4094617
Sequence 15525	Al356534	EST	4108155
Sequence 15526	Al361046	EST	4112667
Sequence 15527	Al361513	EST	4113134
Sequence 15528	Al362439	EST	4114060
Sequence 15529	Al366870	EST	4136615
Sequence 15530			4137677
Sequence 15531	Al367932	EST	
Sequence 15532	A1368934	EST	4147687
Sequence 15533	A1369035	EST	4147788
Sequence 15534	AI371024	EST	4149777
Sequence 15535	Al371230	EST	4149983
Sequence 15536	AI375148	EST	4175138
Sequence 15537	A1375903	EST	4175893
Sequence 15538	Al376130	EST	4176120
Sequence 15539	A1376380	EST	4186229
Sequence 15540	Al376921	EST	4186774
Sequence 15541	AI377037	EST	4186890

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<u>Table 3-5</u>

Sequence 15598	AI565340	EST	4523797
Sequence 15599	Al566010	EST	4524462
Sequence 15600	AI566802	EST	4525254
Sequence 15601	AI567133	EST	4525585
Sequence 15602	AI567453	EST	4525905
Sequence 15603	Al567945	EST	4526397
Sequence 15604	AI569819	EST	4533193
Sequence 15605	AI569945	EST	
Sequence 15606	AI571329	EST	4533319
Sequence 15607	AI571458		4534703
Sequence 15608	AI581200	EST	4534832
Sequence 15609	AI582529	EST	4565576
	AI582700	EST	4568426
Sequence 15610 Sequence 15611	AI597662	EST	4568597
		EST	4606710
Sequence 15612	AI610362	EST	4619529
Sequence 15613	AI619754	EST	4628880
Sequence 15614	Al620259	EST	4629385
Sequence 15615	Al621173	EST	4630299
Sequence 15616	A1624454	EST	4649385
Sequence 15617	AI628617	EST	4665417
Sequence 15618	A1632017	EST	4683347
Sequence 15619	A1632109	EST	4683439
Sequence 15620	A1633086	EST	4684416
Sequence 15621	AI633612	EST	4684942
Sequence 15622	AI634 <b>293</b>	EST	4685623
Sequence 15623	A1635449	EST	4686779
Sequence 15624	AI636182	EST	4687512
Sequence 15625	AI636818	EST	4688148
Sequence 15626	A1638334	EST	4690568
Sequence 15627	AI6507 <b>36</b>	EST	4734715
Sequence 15628 Sequence 15629	AI651228 AI651423	EST	4735207
Sequence 15630	Al651529	EST EST	4735402
Sequence 15631	AI652336	EST	4735508 4736315
Sequence 15632	AI653693	EST	4737672
Sequence 15633	Al653819	EST	4737798
Sequence 15634	AI654736	EST	4738715
Sequence 15635	AI655590	EST	4739569
Sequence 15636	AI655840	EST	4739819
Sequence 15637	AI659386	EST	4762 <b>956</b>
Sequence 15638	AI660901	EST	4764484
Sequence 15639	AI671089	EST	4850820
Sequence 15640	Al673189	EST	4852920
Sequence 15641	Al674314	EST	4874794
Sequence 15642	A1674456	EST	4874936
Sequence 15643	AI674918	EST	4875398
Sequence 15644	AI677810	EST	4887992
Sequence 15645	Al677876	EST	4888058
Sequence 15646	AI679 <b>799</b>	EST	4889981
Sequence 15647	AI680555	EST	4890737
Sequence 15648	A1680742	EST	4890924
Sequence 15649	Al681316	EST	4891498
Sequence 15650	AI683072	EST	4893254
Sequence 15651	Al683117	EST	4893299
Sequence 15652	AI683727	EST	4893909
Sequence 15653	Al689427	EST	4900721
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<u>Table 3-5</u>

Sequence 15710	Al961563	EST	5754276
Sequence 15711	AI971397	EST	5768223
Sequence 15712	Al986177	EST	5813454
Sequence 15713	AL036349	EST	5405930
Sequence 15714	AL037069	EST	5928030
Sequence 15715	AL037845	EST	5407176
		EST	
Sequence 15716	AL037978		5407290
Sequence 15717	AL038564	EST	54 <b>077</b> 50
Sequence 15718	AL039253	EST	5408327
Sequence 15719	AL040602	EST	5409549
Sequence 15720	AL042384	EST	5421723
Sequence 15721	AL046811	EST	5434873
Sequence 15722	AL049135	EST	4728445
Sequence 15723	AL118571	EST	5924470
Sequence 15724	AL120354	EST	5926253
Sequence 15725	AL133982	EST	6602169
Sequence 15726	AL134985	EST	6603172
Sequence 15727	AL138451	EST	6855132
Sequence 15728	AL157600	EST	7058001
Sequence 15729	AW008029	EST	5856807
Sequence 15730	AW008673	EST	5857451
Sequence 15731	AW009152	EST	5857930
Sequence 15732	AW016461	EST	5865218
Sequence 15733	AW020561	EST	5874091
Sequence 15734	AW020914	EST	5874444
Sequence 15735	AW021803	EST	5875333
Sequence 15736	AW022479	EST	5876009
Sequence 15737	AW023580	EST	5877110
Sequence 15738	AW026439	EST	5879969
Sequence 15739	AW026700	EST	5880153
Sequence 15740	AW027717	EST	5886473
Sequence 15741	AW029611	EST	5888367
Sequence 15742	AW051380	EST	5913650
Sequence 15743	AW055331	EST	5921034
Sequence 15744	AW057525	EST	5933164
Sequence 15745	AW062700	EST	6014085
Sequence 15746	AW069166	EST	6024252
Sequence 15747	AW075351	EST	6030349
Sequence 15748	AW079672	EST	6034824
Sequence 15749	AW081110	EST	6036262
Sequence 15750	AW081275	EST	6036427
		EST	6039029
Sequence 15751	AW083877		
Sequence 15752	AW084353	EST	6039505
Sequence 15753	AW084441	EST	6039593
Sequence 15754	AW085581	EST	6040733
Sequence 15755	AW088099	EST	6043904
Sequence 15756	AW089689	EST	6047033
Sequence 15757	AW102924	EST	6073539
1	AW131444	EST	6133051
Sequence 15758			
Sequence 15759	AW151003	EST	6198901
Sequence 15760	AW1511 <b>70</b>	EST	6199068
Sequence 15761	AW161328	EST	6300361
Sequence 15762	AW161668	EST	6300701
Sequence 15763	AW166819	EST	6398344
Sequence 15764	AW167129	EST	6398654
Sequence 15765	AW168050	EST	6399575
Sednetice 19192	V44 1 00000	ESI	0093013

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Sequence 15766	AW177001	EST	6443038
Sequence 15767	AW183074	EST	6451536
Sequence 15768	AW188081	EST	6462517
Sequence 15769	AW188641	EST	6463001
Sequence 15770	AW188668	EST	6463028
Sequence 15771	AW 188775	EST	6463211
Sequence 15772	AW188814	EST	6463250
Sequence 15773	AW 190568	EST	6465048
Sequence 15774	AW235964	EST	6568353
Sequence 15775	AW243325	EST	6577165
Sequence 15776	AW243940	EST	6577780
Sequence 15777	AW271299	EST	6658329
Sequence 15778	AW271852	EST	6658882
Sequence 15779	AW273750	EST	6660780
Sequence 15780	AW290969	EST	6697605
Sequence 15781	AW293305	EST	6699867
Sequence 15782	AW293616	EST	6700252
Sequence 15783	AW294158	EST	6700794
Sequence 15784	AW297885	EST	6704510
Sequence 15785	AW328141	EST	6798637
Sequence 15786	AW360820	EST	6865470
Sequence 15787	AW361122	EST	6865772
Sequence 15788	AW365134	EST	6869784
Sequence 15789	AW368376	EST	6873026
Sequence 15790	AW369111	EST	6873761
Sequence 15791	AW378964	EST	6883623
Sequence 15792	AW381285	EST	6885944
Sequence 15793	AW386920	EST	6891579
Sequence 15794	AW390476	EST	6895135
Sequence 15795	AW394098	EST	6898757
Sequence 15796	AW402596	EST	6921298
Sequence 15797	AW4498 <b>08</b>	EST	6990584
Sequence 15798	AW449890	EST	6990666
Sequence 15799	AW450105	EST	6990881
Sequence 15800	AW 450574	EST	6991350
Sequence 15801	AW450748	EST	6991524
Sequence 15802	AW451232	EST	6992008
Sequence 15803	AW451555	EST	6992331
Sequence 15804	AW451783	EST	6992559
Sequence 15805	AW452088	EST	6992864
Sequence 15806	AW 452730	EST	6993506
Sequence 15807	AW 468168	EST	7038274
Sequence 15808	AW469281	EST	7039387
Sequence 15809	AW470264	EST	7040370
Sequence 15810	AW473121	EST	7043227
Sequence 15811	AW510526	EST	N/A
Sequence 15812	AW511443	EST	N/A
Sequence 15813	AW 572897	EST	7237630
Sequence 15814	C00019	EST	1432249
Sequence 15815	C14194	EST	1568901
Sequence 15816	C16523	EST	1571230
Sequence 15817	D534 <b>8</b> 4	EST	955381
Sequence 15818	D61456	EST	970232
Sequence 15819	D61516	EST	970472
Sequence 15820	D81678	EST	1179555
Sequence 15821	FC8667	EST	672530

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**Table 3-5** 

Sequence 1582	2 F10548	EST	683206
Sequence 1582	3 F35428	EST	4821054
Sequence 1582		EST	885333
Sequence 1582		EST	890789
Sequence 1582		EST	890933
Sequence 1582		EST	915958
Sequence 1582		EST	923470
Sequence 1582		EST	992113
Sequence 1583		EST	992215
		EST	994387
Sequence 1583		EST	1018935
Sequence 1583		EST	1026112
Sequence 1583		EST	1026234
Sequence 1583		EST	1055470
Sequence 1583			1056977
Sequence 1583		EST	
Sequence 1583	7 H79965	EST	1058054
Sequence 1583		EST	1059429
Sequence 1583		EST	1080557
Sequence 1584		EST	1140302
Sequence 1584		EST	1166756
Sequence 1584		EST	1190084
Sequence 1584		EST	1200767
Sequence 1584	4 N62671	EST	1210500
Sequence 1584		EST	1219034
Sequence 1584		EST	1228539
Sequence 1584		EST	1267942
Sequence 1584	8 N95 <b>80</b> 3	EST	1268098
Sequence 1584		EST	778453
Sequence 1585	0 R80268	EST	856549
Sequence 1585		EST	96 <b>7966</b>
Sequence 1585	2 R98534	EST	985051
Sequence 1585		EST	651072
Sequence 1585		EST	657372
Sequence 1585		EST	692046
Sequence 1585		EST	704195
Sequence 1585	7 T84643	EST	712995
Sequence 1585		EST	722981
Sequence 1585	i9 U69 <b>56</b> 7	EST	2731398
Sequence 1586	60 W024 <b>9</b> 0	EST	1274488
Sequence 1586		EST	1276162
Sequence 1586	2 W05733	EST	1278455
Sequence 1586	3 W19899	EST	1295768
Sequence 1586	34 W21 <b>207</b>	EST	1298239
Sequence 1586		EST	1306147
Sequence 1586	6 W58768	EST	1364280
Sequence 1586	7 W60245	EST	1367175
Sequence 1586	8 W61046	EST	1367806
Sequence 1586		EST	1390063
Sequence 1587		EST	288765
Sequence 1587		EST	566739
Sequence 1587		NUCPATENT	N/A
Sequence 1587		NUCPATENT	N/A
Sequence 1587		NUCPATENT	N/A
Sequence 1587		NUCPATENT	N/A
Sequence 158		NUCPATENT	N/A
Sequence 158		NUCPATENT	N/A
_ = = = = = = = = = = = = = = = = = = =			

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<u>Table 3-5</u>

Sequence	15878	V5 <b>8586</b>	NUCPATENT	N/A
Sequence	15879	V5 <b>9067</b>	NUCPATENT	N/A
Sequence	15880	V59630	NUCPATENT	N/A
Sequence	15 <b>881</b>	V60015	NUCPATENT	N/A
Sequence	15882	V65326	NUCPATENT	N/A
Sequence	15883	V88598	NUCPATENT	N/A
Sequence	15884	V90354	NUCPATENT	N/A
Sequence	15885	X02974	NUCPATENT	N/A
Sequence	15886	X20483	NUCPATENT	N/A
Sequence	15887	X26850	NUCPATENT	N/A
Sequence	15888	X27348	NUCPATENT	N/A
Sequence	15889	X30410	NUCPATENT	N/A
Sequence	15890	X33813	NUCPATENT	N/A
Sequence	15891	X84945	NUCPATENT	N/A
Sequence		X86772	NUCPATENT	N/A
Sequence	15893	X87396	NUCPATENT	N/A
Sequence	15894	X87 <b>868</b>	NUCPATENT	N/A
Sequence	15895	X98006	NUCPATENT	N/A
Sequence	158 <b>96</b>	X98010	NUCPATENT	N/A
Sequence	15897	X98409	NUCPATENT	N/A
Sequence	15898	X98714	NUCPATENT	N/A
Sequence:	15899	Z14927	NUCPATENT	N/A
Sequence:	15900	Z16190	NUCPATENT	N/A
Sequence:	15901	Z18356	NUCPATENT	N/A
Secuence:	15902	Z24811	NUCPATENT	N/A
Secuence	15903	Z29718	NUCPATENT	N/A
Sequence:	15904	Z33656	NUCPATENT	N/A
Sequence:	15905	Z34089	NUCPATENT	N/A
Sequence:		Z41384	NUCPATENT	N/A
Sequence :	15907	Z97274	NUCPATENT	N/A
Sequence:	15908	AC28082	PREPATNUC	N/A
Sequence :		AC34864	PREPATNUC	N/A
Sequence :		AC36253	PREPATNUC	N/A
Sequence:	15911	AC38032	PREPATNUC	N/A

PCT/US01/02318 WO 01/53836

## Table 3-6

Sequence 11676: Found in patent publication WO00/05367 AGTACTTCTAGAATTAATTAAGGCGGGCGCCTGCANTCCCAGNTACNCGGNAGGCTGAGG CAGGAGAATGGCGTGAACCCANGAGGCGGAGCTTGCAGTGAGCTGAGATCGCGCCACTGC NCTCCAGCCTGGGCAACAGAGTGAGACTCCGTCCCGCTAANAAATNANAAAANAANAANN CCNGGCCCNCCCNCTGGNGNNTTCCCANATCCCNNTTTTNTGAATTTTTTNCCCCCCCC NNNTGGNCCTGTTTTCTACTAGTGATGATCTGGTAATATACAATTTGTCCAGTAGCCAGT TTGTTTTTATTGTGTTTTCTAACCATAAGAGATCATTAAAGGCAAAGCCTGTATGACGCT **AATCTTCTGGTCA** 

Sequence 11677: Found in patent publication WO99/46375

AGTACTTCTAGAATTAATTAAGGCGGGGACACCACTTTTCAAAGGACTTCTTGGTTTCAG CATAACCTAAGACAGGGAATTGGGAGCCATCATATGTCACAGTGTTCAGAATTCAAGCAT TACATTTAGCTGACAAAGTGCTTATAGAATTTCTTAACAACTGAACCATTCAAAAGGATT TCCATAGCTTCTTATCCCTGTATTGAGGTAATAAATTGTTTTACTGACAATTTTTCCTTT TTCTACACTAAAACAATATGTGATATATTTCCCCTCTTGAAGAGGCAATTCATTAAACTC Sequence 11678: Found in patent publication WO98/54206

ΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑ

Sequence 11679: Found in patent publication WO99/63088

AGTACTTCTAGAATTAATTAAGGCGGGGAGTTTTATCTATTCTTTACATTGGATGCAAAA TGTATTACTCAAGAAGAGGCATTCGGTATCGAACCATAGATGAACATGATGCCATCATTT GTTTATTAATAGTTTAAAACAATATTCTCTTTTTGAAAATAGTATAAACAGGCCATGCAT ATAATGTACAGTGTATTACGTAAATATGTAAAGATTCTTCAAGGTAACAAGGGTTTGGGT TTTGAAATAAACATCTGGATCTTATAGACCGTTCATACAATGGTTTTAGCAAGTTCATAG TAAGACAAACAAGTCCTATCTTTTTTTTTTTTTGGCTGGGGTGGGGGCATTGGTCACATAT **GACCAGTA** 

Sequence 11680: Found in patent publication WO99/67383

AGTACTTCTAGAATTAATTAAGGCGGGTCCCTGCTTGTCCTATGAACTGCTCAGAGCTCC TGTCAGTCCAGCTGGGCCTTCTGGGTTCTGGCACCATTTCGTAGCCATTCTCTTTGTATT TTAAAAGGACGTTATGAAAGGGCTTAGACCAAAATAAATCATAATGTTACTTGAGCCACC 

Sequence 13322: Found in patent publication WO00/09552

ΑΛΑΛΑΛΑΛΑΛΑΛΑΛΑΛΑΛΑΛΑΛΑΛΑΛΑΛΑΛΑΛΑΛΑΛΑ

Sequence 13323: Found in patent publication WO99/66041

AGTACTTCTAGAATTAATTAAGGCGGGGGAAAGTTAGAGGAACTGAAAGTTTGGGAATAG GCTGACCACATATTATGCCAGTGACCAGTATGACAGGAGATGGGGCCCTGCTGCCAGTCA 

Sequence 13324: Found in patent publication WO99/66041

Sequence 13325: Found in patent publication WO99/67383

AGTACTTCTAGAATTAATTAAGGCGGGTCCCTGCTTGTCCTATGAACTGCTCAGAGCTCC TGTCAGTCCAGCTGGGCCTTCTGGGTTCTGGCACCATTTCGTAGCCATTCTCTTTGTATT TTAAAAGGACGTTATGAAAGGGCTTAGACCAAAATAAATCATAATGTTACTTGAGCCACC TTATATAGCTGCTTGGAGAGTCTGTGTAGTTCTTTCTGCATGTATTAAAAATGTTTAGAA

Sequence 13326: Found in patent publication WO00/06698

GCCCTGGGAAACGATTGTGAACGCNTGAATGAATTGATGACTAANATCCGCTGCGGGGGT CCTACAGCGCANATGGTAATGCCCGTTCTGACTGGCTGGGAACGGCACCTTAGCAAGATA CTTAAAAGGCGCCTTCTGTGTGCCACNTGCACTCCACCCTGGGCAACAGAGCAAANACCC

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### **Table 3-6**

CATCTCAAAAATAAATAAATATATATAAAAAAATAAAAAGCTATTTCTAGTTTNATTTCAC TATAAAGTTTTGCTTTTATTAAA Sequence 13327: Found in patent publication US6034229 AGTACTTCTAGAATTAATTAAGGCGGGCGGATCATGAGGTCAGGAGTTCGAGACCAGCCT GGCCAACATGGTGAAACTCCGTGTCTACTAAAAATACAAAAATTGGCTGGGCCGTGGTGG TTCGCACCTGTAGTCCCAGCTACTGGGGGGGGCTGAGGCAGGAGAATGGCTTGAACCCGGG AGGGTAGAGGTTGCAGTGAGCCAAGATTGTGCCACTGCACTCCAGCCTGGGCGACTGAGC AAGACTGTCTAAAAAAAAAAAAAAAAAAAATAAATAAAGNAA Sequence 13328: Found in patent publication WO00/18925 GCCGTCGTTTTACACCCCGNAANAANCNATANCNATGATNGNTGCCNGTCCCNCTCNNNA NAGNAATAAAAANTAATTAANGCGGGTGATTAAATTTCAGTCCATTATGATTTTCCTT TCTCACATAATTACTTTTTCTTTTTAGACTTATAAGCTAGCAATTACAGATTTAACTAC AGCTATCAGCATGGACAAAAATAGTTATACAGCATTTTATAACAGAGCATTATGTTACAC CAAGATAAGGGAACTTCAAATGGCATTAACAGATTATGGAATTGTGCTGCTTCTTGATGC TACAGAAACTGTNAAACTAAATACCTTCCTTAATCGTGGACTCATCTACGTAGAACTAGG CCAGTATGGCTTTGCACTAGAGGATTTTAAAAC Sequence 15908: Found in patent publication WO98/33916 AAATTNNNTATTTATTTTAAANTAAAAANTATACTNAACTAATTATTTNAATTAAAGAAA Sequence 15909: Found in patent publication WO99/14328 Sequence 15910: Found in patent publication WO99/66041 Sequence 15911: Found in patent publication WO00/05382 AAGACATTCCAGACAAAAAGACTAACTTGTCAGAAAGCCCTGTGGCGGAAGGGAGCTTTT TTTGTAAAGGAGTTTTGGTTTTCTCCTAATAGCAATGGGATATCTTCCAAGGAATCTCAA TCAAAAGGGAGAGATGGCTCCGATTGGAATGTCATCCCTGGCTGAAGAGTAGAGGAAGCA AAAAAAAAAAAAAAAAAAANNGNNNNNNCCCNNTNTTTTTTT

## Table 3-6

TTCTACACTAAAACAATATGTGATATATTTCCCCTCTTGAAGAGGCAATTCATTAAACTC Sequence 11678: Found in patent publication WO98/54206

Sequence 11679: Found in patent publication WO99/63088

Sequence 11680: Found in patent publication WO99/67383

Sequence 13322: Found in patent publication WO00/09552

Sequence 13323: Found in patent publication WO99/66041

Sequence 13324: Found in patent publication WO99/66041

Sequence 13325: Found in patent publication WO99/67383

AGTACTTCTAGAATTAATTAAGGCGGGTCCCTGCTTGTCCTATGAACTGCTCAGAGCTCC TGTCAGTCCAGCTGGGCCTTCTGGGTTCTGGCACCATTTCGTAGCCATTCTCTTTGTATT TTAAAAGGACGTTATGAAAGGGCTTAGACCAAAATAAATCATAATGTTACTTGAGCCACC TTATATAGCTGCTTGGAGAGTCTGTGTAGTTCTTTCTGCATGTATTAAAAATGTTTAGAA

Sequence 13326: Found in patent publication WO00/06698

GCCTGGGAAACGATTGTGAACGCNTGAATGAATTGATGACTAANATCCGCTGCGGGGGT CCTACAGCGCANATGGTAATGCCCGTTCTGACTGGCTGGGAACGCACCTTAGCAAGATA CTTAAAAGGCGCCTTCTGTGTGCCACNTGCACTCCACCCTGGGCAACAGAGCAAANACCC

#### Table 3-6

CATCTCAAAAATAAATAAATATATATAAAAAAATAAAAAGCTATTTCTAGTTTNATTTCAC TATAAAGTTTTGCTTTTATTAAA Sequence 13327: Found in patent publication US6034229 AGTACTTCTAGAATTAATTAAGGCGGCGGATCATGAGGTCAGGAGTTCGAGACCAGCCT GGCCAACATGGTGAAACTCCGTGTCTACTAAAAATACAAAAATTGGCTGGGCCGTGGTGG TTCGCACCTGTAGTCCCAGCTACTGGGGGGGGCTGAGGCAGGAGAATGGCTTGAACCCGGG AGGGTAGAGGTTGCAGTGAGCCAAGATTGTGCCACTGCACTCCAGCCTGGGCGACTGAGC AAGACTGTCTAAAAAAAAAAAAAAAAATAAATAAAGNAA Sequence 13328: Found in patent publication WO00/18925 GCCGTCGTTTTACACCCCGNAANAANCNATANCNATGATNGNTGCCNGTCCCCCTCNNNA NAGNAATAAAAANTAATTAANGCGGGTGATTAAATTTCAGTCCATTATGATTTTCCTT TCTCACATAATTACTTTTTCTTTTTAGACTTATAAGCTAGCAATTACAGATTTAACTAC AGCTATCAGCATGGACAAAAATAGTTATACAGCATTTTATAACAGAGCATTATGTTACAC CAAGATAAGGGAACTTCAAATGGCATTAACAGATTATGGAATTGTGCTGCTTCTTGATGC TACAGAAACTGTNAAACTAAATACCTTCCTTAATCGTGGACTCATCTACGTAGAACTAGG CCAGTATGGCTTTGCACTAGAGGATTTTAAAAC Sequence 15908: Found in patent publication WO98/33916 AAATTNNNTATTTAATTTAAANTAAAAANTATACTNAACTAATTATTTNAATTAAAGAAA Sequence 15909: Found in patent publication WO99/14328 Sequence 15910: Found in patent publication WO99/66041 Sequence 15911: Found in patent publication WO00/05382 AAGACATTCCAGACAAAAAGACTAACTTGTCAGAAAGCCCTGTGGCGGAAGGGAGCTTTT TTTGTAAAGGAGTTTTGGTTTTCTCCTAATAGCAATGGGATATCTTCCAAGGAATCTCAA TCAAAAGGGAGAGATGGCTCCGATTGGAATGTCATCCCTGGCTGAAGAGTAGAGGAAGCA AAAAAAAAAAAAAAAAAAAANNGNNNNNNCCCNNTNTTTTTTT

#### Table 3-6

Sequence 11677: Found in patent publication WO99/46375

Sequence 11679: Found in patent publication WO99/63088

Sequence 11680: Found in patent publication WO99/67383

Sequence 13322: Found in patent publication WO00/09552

Sequence 13323: Found in patent publication WO99/66041

Sequence 13324: Found in patent publication WO99/66041

Sequence 13325: Found in patent publication WO99/67383

AGTACTTCTAGAATTAATTAAGGCGGGTCCCTGCTTGTCCTATGAACTGCTCAGAGCTCC TGTCAGTCCAGCTGGGCCTTCTGGGTTCTGGCACCATTTCGTAGCCATTCTCTTTGTATT TTAAAAGGACGTTATGAAAGGGCTTAGACCAAAATAAATCATAATGTTACTTGAGCCACC TTATATAGCTGCTTGGAGAGTCTGTGTAGTTCTTTCTGCATGTATTAAAAATGTTTAGAA

Sequence 13326: Found in patent publication WO00/06698

GCCTGGGAAACGATTGTGAACGCNTGAATGAATTGATGACTAANATCCGCTGCGGGGGT CCTACAGCGCANATGGTAATGCCCGTTCTGACTGGCTGGGAACGGCACCTTAGCAAGATA CTTAAAAGGCGCCTTCTGTGTGCCACNTGCACTCCACCCTGGGCAACAGAGCAAANACCC

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#### Table 3-6

CATCTCAAAAATAAATAAATATATATAAAAAAATAAAAAGCTATTTCTAGTTTNATTTCAC TATAAAGTTTTGCTTTTATTAAA Sequence 13327: Found in patent publication US6034229 AGTACTTCTAGAATTAATTAAGGCGGCGGATCATGAGGTCAGGAGTTCGAGACCAGCCT GGCCAACATGGTGAAACTCCGTGTCTACTAAAAATACAAAAATTGGCTGGGCCGTGGTGG AGGGTAGAGGTTGCAGTGAGCCAAGATTGTGCCACTGCACTCCAGCCTGGGCGACTGAGC AAGACTGTCTAAAAAAAAAAAAAAAATAAATAAAGNAA Sequence 13328: Found in patent publication WO00/18925 GCCGTCGTTTTACACCCCGNAANAANCNATANCNATGATNGNTGCCNGTCCCNCTCNNNA NAGNAATAAAAANTAATTAANGCGGGTGATTAAATTTCAGTCCATTATGATTTTCCTT TCTCACATAATTACTTTTTTTTTTTTTAGACTTATAAGCTAGCAATTACAGATTTAACTAC AGCTATCAGCATGGACAAAAATAGTTATACAGCATTTTATAACAGAGCATTATGTTACAC CAAGATAAGGGAACTTCAAATGGCATTAACAGATTATGGAATTGTGCTGCTTCTTGATGC TACAGAAACTGTNAAACTAAATACCTTCCTTAATCGTGGACTCATCTACGTAGAACTAGG CCAGTATGGCTTTGCACTAGAGGATTTTAAAAC Sequence 15908: Found in patent publication WO98/33916 AAATTNNNTATTTATTTTAAANTAAAAANTATACTNAACTAATTATTTNAATTAAAGAAA AAATAANTAAAAAAATTNATAAANAATAAATNTTA AATAAAAAATATT Sequence 15909: Found in patent publication WO99/14328 Sequence 15910: Found in patent publication WO99/66041 Sequence 15911: Found in patent publication WO00/05382 AAGACATTCCAGACAAAAAGACTAACTTGTCAGAAAGCCCTGTGGCGGAAGGGAGCTTTT TTTGTAAAGGAGTTTTGGTTTTCTCCTAATAGCAATGGGATATCTTCCAAGGAATCTCAA TCAAAAGGGAGAGATGGCTCCGATTGGAATGTCATCCCTGGCTGAAGAGTAGAGGAAGCA AAAAAAAAAAAAAAAAAAAANNGNNNNNNCCCNNTNTTTTTTT

#### **Table 3-6**

Sequence 11676: Found in patent publication WO00/05367 AGTACTTCTAGAATTAATTAAGGCGGGCGCCTGCANTCCCAGNTACNCGGNAGGCTGAGG CAGGAGAATGGCGTGAACCCANGAGGCGGAGCTTGCAGTGAGCTGAGATCGCGCCACTGC NCTCCAGCCTGGGCAACAGAGTGAGACTCCGTCCCGCTAANAAATNANAAAANAANAANN CONGCCONCCONCTGGNGNNTTCCCANATCCCNNTTTTNTGAATTTTTTNCCCCCCCC NNNTGGNCCTGTTTTCTACTAGTGATGATCTGGTAATATACAATTTGTCCAGTAGCCAGT TTGTTTTTATTGTGTTTTCTAACCATAAGAGATCATTAAAGGCAAAGCCTGTATGACGCT AATCTTCTGGTCA Sequence 11677: Found in patent publication WO99/46375 AGTACTTCTAGAATTAATTAAGGCGGGGACACCACTTTTCAAAGGACTTCTTGGTTTCAG CATAACCTAAGACAGGGAATTGGGAGCCATCATATGTCACAGTGTTCAGAATTCAAGCAT TACATTTAGCTGACAAAGTGCTTATAGAATTTCTTAACAACTGAACCATTCAAAAGGATT TCCATAGCTTCTTATCCCTGTATTGAGGTAATAAATTGTTTTACTGACAATTTTTCCTTT TTCTACACTAAAACAATATGTGATATATTTCCCCTCTTGAAGAGGCAATTCATTAAACTC Sequence 11678: Found in patent publication WO98/54206 AAAAAAAAAAAAAAAAAAAAAAAAAAA Sequence 11679: Found in patent publication WO99/63088 AGTACTTCTAGAATTAATTAAGGCGGGGGGTTTTATCTATTCTTTACATTGGATGCAAAA TGTATTACTCAAGAAGAGGCATTCGGTATCGAACCATAGATGAACATGATGCCATCATTT GTTTATTAATAGTTTAAAACAATATTCTCTTTTTGAAAAATAGTATAAACAGGCCATGCAT ATAATGTACAGTGTATTACGTAAATATGTAAAGATTCTTCAAGGTAACAAGGGTTTGGGT TTTGAAATAAACATCTGGATCTTATAGACCGTTCATACAATGGTTTTAGCAAGTTCATAG TAAGACAAACAAGTCCTATCTTTTTTTTTTTTGGCTGGGGTGGGGGCATTGGTCACATAT **GACCAGTA** Sequence 11680: Found in patent publication WO99/67383 AGTACTTCTAGAATTAATTAAGGCGGGTCCCTGCTTGTCCTATGAACTGCTCAGAGCTCC TGTCAGTCCAGCTGGGCCTTCTGGGTTCTGGCACCATTTCGTAGCCATTCTCTTTGTATT TTAAAAGGACGTTATGAAAGGGCTTAGACCAAAATAAATCATAATGTTACTTGAGCCACC Sequence 13322: Found in patent publication WO00/09552 ΑΑΑΑΑΑΑΑΝΤΤΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑ Sequence 13323: Found in patent publication WO99/66041 AGTACTTCTAGAATTAATTAAGGCGGGGGAAAGTTAGAGGAACTGAAAGTTTGGGAATAG GCTGACCACATATTATGCCAGTGACCAGTATGACAGGAGATGGGGCCCTGCTGCCAGTCA Sequence 13324: Found in patent publication WO99/66041 Sequence 13325: Found in patent publication WO99/67383 AGTACTTCTAGAATTAATTAAGGCGGGTCCCTGCTTGTCCTATGAACTGCTCAGAGCTCC TGTCAGTCCAGCTGGGCCTTCTGGGTTCTGGCACCATTTCGTAGCCATTCTCTTTGTATT TTAAAAGGACGTTATGAAAGGGCTTAGACCAAAATAAATCATAATGTTACTTGAGCCACC TTATATAGCTGCTTGGAGAGTCTGTGTAGTTCTTTCTGCATGTATTAAAAATGTTTAGAA Sequence 13326: Found in patent publication WO00/06698 GCCCTGGGAAACGATTGTGAACGCNTGAATGAATTGATGACTAANATCCGCTGCGGGGGT CCTACAGCGCANATGGTAATGCCCGTTCTGACTGGCTGGGAACGGCACCTTAGCAAGATA

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CTTAAAAGGCGCCTTCTGTGTGCCACNTGCACTCCACCCTGGGCAACAGAGCAAANACCC

#### **Table 3-6**

CATCTCAAAAATAAATAAATATATATAAAAAAATAAAAAGCTATTTCTAGTTTNATTTCAC TATAAAGTTTTGCTTTTATTAAA Sequence 13327: Found in patent publication US6034229 AGTACTTCTAGAATTAATTAAGGCGGGCGGATCATGAGGTCAGGAGTTCGAGACCAGCCT GGCCAACATGGTGAAACTCCGTGTCTACTAAAAATACAAAAATTGGCTGGGCCGTGGTGG TTCGCACCTGTAGTCCCAGCTACTGGGGGGGGGCTGAGGCAGGAGAATGGCTTGAACCCGGG AGGGTAGAGGTTGCAGTGAGCCAAGATTGTGCCACTGCACTCCAGCCTGGGCGACTGAGC AAGACTGTCTAAAAAAAAAAAAAAAAAAATAAATAAAGNAA Sequence 13328: Found in patent publication WO00/18925 GCCGTCGTTTTACACCCCGNAANAANCNATANCNATGATNGNTGCCNGTCCCNCTCNNNA NAGNAATAATAAAANTAATTAANGCGGGTGATTAAATTTCAGTCCATTATGATTTTCCTT TCTCACATAATTACTTTTTCTTTTTAGACTTATAAGCTAGCAATTACAGATTTAACTAC AGCTATCAGCATGGACAAAAATAGTTATACAGCATTTTATAACAGAGCATTATGTTACAC CAAGATAAGGGAACTTCAAATGGCATTAACAGATTATGGAATTGTGCTGCTTCTTGATGC TACAGAAACTGTNAAACTAAATACCTTCCTTAATCGTGGACTCATCTACGTAGAACTAGG CCAGTATGGCTTTGCACTAGAGGATTTTAAAAC Sequence 15908: Found in patent publication WO98/33916 AACAACAAAAAAAACNAAAAATANAAAAAAAAAAAAAANAANNANNCAANNAAAANTAA AAATTNNNTATTTATTTTAAANTAAAAANTATACTNAACTAATTATTTNAATTAAAGAAA Sequence 15909: Found in patent publication WO99/14328 Sequence 15910: Found in patent publication WO99/66041 Sequence 15911: Found in patent publication WO00/05382 AAGACATTCCAGACAAAAAGACTAACTTGTCAGAAAGCCCTGTGGCGGAAGGGAGCTTTT CCAATATGAAGAACTGAGCCTGGAGAGATGGGGATGAGGGGGGAGTGTCGAACCTTTTAGGC TTTGTAAAGGAGTTTTGGTTTTCTCCTAATAGCAATGGGATATCTTCCAAGGAATCTCAA TCAAAAGGGAGAGATGGCTCCGATTGGAATGTCATCCCTGGCTGAAGAGTAGAGGAAGCA AAAAAAAAAAAAAAAAAAAAANNGNNNNNNCCCNNTNTTTTTTT

PCT/US01/02318 WO 01/53836

#### **Table 3-6**

Sequence 11676: Found in patent publication WO00/05367 AGTACTTCTAGAATTAATTAAGGCGGGCGCCTGCANTCCCAGNTACNCGGNAGGCTGAGG CAGGAGAATGGCGTGAACCCANGAGGCGGAGCTTGCAGTGAGCTGAGATCGCGCCACTGC NCTCCAGCCTGGGCAACAGAGTGAGACTCCGTCCCGCTAANAAATNANAAAANAANAANN CCNGGCCCNCCCNCTGGNGNNTTCCCANATCCCNNTTTTNTGAATTTTTTNCCCCCCCC NNNTGGNCCTGTTTTCTACTAGTGATGATCTGGTAATATACAATTTGTCCAGTAGCCAGT TTGTTTTTATTGTGTTTTCTAACCATAAGAGATCATTAAAGGCAAAGCCTGTATGACGCT **AATCTTCTGGTCA** 

Sequence 11677: Found in patent publication WO99/46375

AGTACTTCTAGAATTAATTAAGGCGGGGACACCACTTTTCAAAGGACTTCTTGGTTTCAG CATAACCTAAGACAGGGAATTGGGAGCCATCATATGTCACAGTGTTCAGAATTCAAGCAT TACATTTAGCTGACAAAGTGCTTATAGAATTTCTTAACAACTGAACCATTCAAAAGGATT TCCATAGCTTCTTATCCCTGTATTGAGGTAATAAATTGTTTTACTGACAATTTTTCCTTT TTCTACACTAAAACAATATGTGATATATTTCCCCTCTTGAAGAGGCAATTCATTAAACTC Sequence 11678: Found in patent publication WO98/54206

Sequence 11679: Found in patent publication WO99/63088

AGTACTTCTAGAATTAATTAAGGCGGGGAGTTTTATCTATTCTTTACATTGGATGCAAAA TGTATTACTCAAGAAGAGGCATTCGGTATCGAACCATAGATGAACATGATGCCATCATTT GTTTATTAATAGTTTAAAACAATATTCTCTTTTTGAAAATAGTATAAACAGGCCATGCAT ATAATGTACAGTGTATTACGTAAATATGTAAAGATTCTTCAAGGTAACAAGGGTTTGGGT TTTGAAATAAACATCTGGATCTTATAGACCGTTCATACAATGGTTTTAGCAAGTTCATAG TAAGACAAACAAGTCCTATCTTTTTTTTTTTTGGCTGGGGTGGGGGCATTGGTCACATAT GACCAGTA

Sequence 11680: Found in patent publication WO99/67383

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Sequence 13322: Found in patent publication WO00/09552

Sequence 13323: Found in patent publication WO99/66041

AGTACTTCTAGAATTAATTAAGGCGGGGGAAAGTTAGAGGAACTGAAAGTTTGGGAATAG GCTGACCACATATTATGCCAGTGACCAGTATGACAGGAGATGGGGCCCTGCTGCCAGTCA 

Sequence 13324: Found in patent publication WO99/66041

Sequence 13325: Found in patent publication WO99/67383

AGTACTTCTAGAATTAATTAAGGCGGGTCCCTGCTTGTCCTATGAACTGCTCAGAGCTCC TGTCAGTCCAGCTGGGCCTTCTGGGTTCTGGCACCATTTCGTAGCCATTCTCTTTGTATT TTAAAAGGACGTTATGAAAGGGCTTAGACCAAAATAAATCATAATGTTACTTGAGCCACC TTATATAGCTGCTTGGAGAGTCTGTGTAGTTCTTTCTGCATGTATTAAAAATGTTTAGAA

Sequence 13326: Found in patent publication WO00/06698 GCCCTGGGAAACGATTGTGAACGCNTGAATGAATTGATGACTAANATCCGCTGCGGGGGT CCTACAGCGCANATGGTAATGCCCGTTCTGACTGGCTGGGAACGGCACCTTAGCAAGATA CTTAAAAGGCGCCTTCTGTGTGCCACNTGCACTCCACCCTGGGCAACAGAGCAAANACCC

#### **Table 3-6**

CATCTCAAAAATAAATAAATATATATAAAAAAATAAAAAGCTATTTCTAGTTTNATTTCAC TATAAAGTTITGCTTTTATTAAA Sequence 13327: Found in patent publication US6034229 AGTACTTCTAGAATTAATTAAGGCGGGCGGATCATGAGGTCAGGAGTTCGAGACCAGCCT GGCCAACATGGTGAAACTCCGTGTCTACTAAAAATACAAAAATTGGCTGGGCCGTGGTGG TTCGCACCTGTAGTCCCAGCTACTGGGGGGGGGCTGAGGCAGGAGAATGGCTTGAACCCGGG AGGGTAGAGGTTGCAGTGAGCCAAGATTGTGCCACTGCACTCCAGCCTGGGCGACTGAGC AAGACTGTCTAAAAAAAAAAAAAAAAAAATAAATAAAGNAA Sequence 13328: Found in patent publication WO00/18925 GCCGTCGTTTTACACCCCGNAANAANCNATANCNATGATNGNTGCCNGTCCCNCTCNNNA NAGNAATAATAAAANTAATTAANGCGGGTGATTAAATTTCAGTCCATTATGATTTTCCTT TCTCACATAATTACTTTTTTCTTTTTAGACTTATAAGCTAGCAATTACAGATTTAACTAC AGCTATCAGCATGGACAAAAATAGTTATACAGCATTTTATAACAGAGCATTATGTTACAC CAAGATAAGGGAACTTCAAATGGCATTAACAGATTATGGAATTGTGCTGCTTCTTGATGC TACAGAAACTGTNAAACTAAATACCTTCCTTAATCGTGGACTCATCTACGTAGAACTAGG CCAGTATGGCTTTGCACTAGAGGATTTTAAAAC Sequence 15908: Found in patent publication WO98/33916 AACAACAAAAAAACNAAAAATANAAAAAAAAAAAAAAANAANNNANNCAANNNAAAANTAA AAATTNNNTATTTATTTTAAANTAAAAANTATACTNAACTAATTATTTNAATTAAAGAAA Sequence 15909: Found in patent publication WO99/14328 ΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑ Sequence 15910: Found in patent publication WO99/66041 Sequence 15911: Found in patent publication WO00/05382 AAGACATTCCAGACAAAAAGACTAACTTGTCAGAAAGCCCTGTGGCGGAAGGGAGCTTTT CCAATATGAAGAACTGAGCCTGGAGAGATGGGATGAGGGGGGAGTGTCGAACCTTTTAGGC TTTGTAAAGGAGTTTTGGTTTTCTCCTAATAGCAATGGGATATCTTCCAAGGAATCTCAA TCAAAAGGGAGAGATGGCTCCGATTGGAATGTCATCCCTGGCTGAAGAGTAGAGGAAGCA AAAAAAAAAAAAAAAAAAAAAAANNGNNNNNNCCCNNTNTTTTTTT

PCT/US01/02318 WO 01/53836

#### Table 3-6

Sequence 11676: Found in patent publication WO00/05367 AGTACTTCTAGAATTAATTAAGGCGGGCGCCTGCANTCCCAGNTACNCGGNAGGCTGAGG CAGGAGAATGGCGTGAACCCANGAGGCGGAGCTTGCAGTGAGCTGAGATCGCGCCACTGC NCTCCAGCCTGGGCAACAGAGTGAGACTCCGTCCCGCTAANAAATNANAAAANAANAANN NNNTGGNCCTGTTTCTACTAGTGATGATCTGGTAATATACAATTTGTCCAGTAGCCAGT TTGTTTTTATTGTGTTTTCTAACCATAAGAGATCATTAAAGGCAAAGCCTGTATGACGCT **AATCTTCTGGTCA** 

Sequence 11677: Found in patent publication WO99/46375

AGTACTTCTAGAATTAATTAAGGCGGGGACACCACTTTTCAAAGGACTTCTTGGTTTCAG CATAACCTAAGACAGGGAATTGGGAGCCATCATATGTCACAGTGTTCAGAATTCAAGCAT TACATTTAGCTGACAAAGTGCTTATAGAATTTCTTAACAACTGAACCATTCAAAAGGATT TTTTTGTTTAAAACTGGATTTCAATGTAAGCAAATGAAGAAAAAAATATAGATTTCATT TCCATAGCTTCTTATCCCTGTATTGAGGTAATAAATTGTTTTACTGACAATTTTTCCTTT TTCTACACTAAAACAATATGTGATATATTTCCCCTCTTGAAGAGGCAATTCATTAAACTC Sequence 11678: Found in patent publication WO98/54206

AAAAAAAAAAAAAAAAAAAAAA

Sequence 11679: Found in patent publication WO99/63088

AGTACTTCTAGAATTAATTAAGGCGGGGGGTTTTATCTATTCTTTACATTGGATGCAAAA TGTATTACTCAAGAAGAGGCATTCGGTATCGAACCATAGATGAACATGATGCCATCATTT GTTTATTAATAGTTTAAAACAATATTCTCTTTTTTGAAAATAGTATAAACAGGCCATGCAT ATAATGTACAGTGTATTACGTAAATATGTAAAGATTCTTCAAGGTAACAAGGGTTTGGGT TITGAAATAAACATCTGGATCTTATAGACCGTTCATACAATGGTTTTAGCAAGTTCATAG TAAGACAAACAAGTCCTATCTTTTTTTTTTTTGGCTGGGGTGGGGGCATTGGTCACATAT **GACCAGTA** 

Sequence 11680: Found in patent publication WO99/67383 AGTACTTCTAGAATTAATTAAGGCGGGTCCCTGCTTGTCCTATGAACTGCTCAGAGCTCC TGTCAGTCCAGCTGGGCCTTCTGGGTTCTGGCACCATTTCGTAGCCATTCTCTTTGTATT

TTAAAAGGACGTTATGAAAGGGCTTAGACCAAAATAAATCATAATGTTACTTGAGCCACC 

Sequence 13322: Found in patent publication WO00/09552

Sequence 13323: Found in patent publication WO99/66041

AGTACTTCTAGAATTAATTAAGGCGGGGGAAAGTTAGAGGAACTGAAAGTTTGGGAATAG GCTGACCACATATTATGCCAGTGACCAGTATGACAGGAGATGGGGCCCTGCTGCCAGTCA 

Sequence 13324: Found in patent publication WO99/66041

Sequence 13325: Found in patent publication WO99/67383

AGTACTTCTAGAATTAATTAAGGCGGGTCCCTGCTTGTCCTATGAACTGCTCAGAGCTCC TGTCAGTCCAGCTGGGCCTTCTGGGTTCTGGCACCATTTCGTAGCCATTCTCTTTGTATT TTAAAAGGACGTTATGAAAGGGCTTAGACCAAAATAAATCATAATGTTACTTGAGCCACC TTATATAGCTGCTTGGAGAGTCTGTGTAGTTCTTTCTGCATGTATTAAAAATGTTTAGAA

Sequence 13326: Found in patent publication WO00/06698

GCCCTGGGAAACGATTGTGAACGCNTGAATGAATTGATGACTAANATCCGCTGCGGGGGT CCTACAGCGCANATGGTAATGCCCGTTCTGACTGGCTGGGAACGGCACCTTAGCAAGATA CTTAAAAGGCGCCTTCTGTGTGCCACNTGCACTCCACCCTGGGCAACAGAGCAAANACCC

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#### Table 3-6

CATCTCAAAAATAAATAAATATATATAAAAAAATAAAAAGCTATTTCTAGTTTNATTTCAC TATAAAGTTTTGCTTTTATTAAA Sequence 13327: Found in patent publication US6034229 AGTACTTCTAGAATTAATTAAGGCGGCGGCGATCATGAGGTCAGGAGTTCGAGACCAGCCT GGCCAACATGGTGAAACTCCGTGTCTACTAAAAATACAAAAATTGGCTGGGCCGTGGTGG TTCGCACCTGTAGTCCCAGCTACTGGGGGGGGCTGAGGCAGGAGAATGGCTTGAACCCGGG AGGGTAGAGGTTGCAGTGAGCCAAGATTGTGCCACTGCACTCCAGCCTGGGCGACTGAGC AAGACTGTCTAAAAAAAAAAAAAAAAATAAATAAAGNAA Sequence 13328: Found in patent publication WO00/18925 GCCGTCGTTTTACACCCCGNAANAANCNATANCNATGATNGNTGCCNGTCCCNCTCNNNA NAGNAATAAAANTAATTAANGCGGGTGATTAAATTTCAGTCCATTATGATTTTCCTT TCTCACATAATTACTTTTTTTTTTTAGACTTATAAGCTAGCAATTACAGATTTAACTAC AGCTATCAGCATGGACAAAAATAGTTATACAGCATTTTATAACAGAGCATTATGTTACAC CAAGATAAGGGAACTTCAAATGGCATTAACAGATTATGGAATTGTGCTGCTTCTTGATGC TACAGAAACTGTNAAACTAAATACCTTCCTTAATCGTGGACTCATCTACGTAGAACTAGG CCAGTATGGCTTTGCACTAGAGGATTTTAAAAC Sequence 15908: Found in patent publication WO98/33916 AAATTNNNTATTTATTTTAAANTAAAAANTATACTNAACTAATTATTTNAATTAAAGAAA Sequence 15909: Found in patent publication WO99/14328 Sequence 15910: Found in patent publication WO99/66041 Sequence 15911: Found in patent publication WO00/05382 AAGACATTCCAGACAAAAAGACTAACTTGTCAGAAAGCCCTGTGGCGGAAGGGAGCTTTT CCAATATGAAGAACTGAGCCTGGAGAGATGGGATGAGGGGGGAGTGTCGAACCTTTTAGGC TTTGTAAAGGAGTTTTGGTTTTCCCTAATAGCAATGGGATATCTTCCAAGGAATCTCAA TCAAAAGGGAGAGATGGCTCCGATTGGAATGTCATCCCTGGCTGAAGAGTAGAGGAAGCA

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PCT/US01/02318 WO 01/53836

#### **Table 3-6**

Sequence 11676: Found in patent publication WC00/05367 AGTACTTCTAGAATTAATTAAGGCGGGCGCCTGCANTCCCAGNTACNCGGNAGGCTGAGG CAGGAGAATGGCGTGAACCCANGAGGCGGAGCTTGCAGTGAGCTGAGATCGCGCCACTGC NCTCCAGCCTGGGCAACAGAGTGAGACTCCGTCCCGCTAANAAATNANAAAANAANAANN CCNGGCCCNCCCNCTGGNGNNTTCCCANATCCCNNTTTTNTGAATTTTTTTNCCCCCCCC NNNTGGNCCTGTTTTCTACTAGTGATGATCTGGTAATATACAATTTGTCCAGTAGCCAGT TTGTTTTTATTGTGTTTTCTAACCATAAGAGATCATTAAAGGCAAAGCCTGTATGACGCT **AATCTTCTGGTCA** 

Sequence 11677: Found in patent publication WO99/46375

AGTACTTCTAGAATTAATTAAGGCGGGGACACCACTTTTCAAAGGACTTCTTGGTTTCAG CATAACCTAAGACAGGGAATTGGGAGCCATCATATGTCACAGTGTTCAGAATTCAAGCAT TACATTTAGCTGACAAAGTGCTTATAGAATTTCTTAACAACTGAACCATTCAAAAGGATT TCCATAGCTTCTTATCCCTGTATTGAGGTAATAAATTGTTTTACTGACAATTTTTCCTTT TTCTACACTAAAACAATATGTGATATTTTCCCCTCTTGAAGAGGCAATTCATTAAACTC Sequence 11678: Found in patent publication WO98/54206

AAAAAAAAAAAAAAAAAAAAAAAA

Sequence 11679: Found in patent publication WO99/63088

AGTACTTCTAGAATTAATTAAGGCGGGGGGTTTTATCTATTCTTTACATTGGATGCAAAA TGTATTACTCAAGAAGAGGCATTCGGTATCGAACCATAGATGAACATGATGCCATCATTT GTTTATTAATAGTTTAAAACAATATTCTCTTTTTGAAAATAGTATAAACAGGCCATGCAT ATAATGTACAGTGTATTACGTAAATATGTAAAGATTCTTCAAGGTAACAAGGGTTTGGGT TTTGAAATAAACATCTGGATCTTATAGACCGTTCATACAATGGTTTTAGCAAGTTCATAG TAAGACAAACAAGTCCTATCTTTTTTTTTTTGGCTGGGGTGGGGGCATTGGTCACATAT GACCAGTA

Sequence 11680: Found in patent publication WO99/67383

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Sequence 13322: Found in patent publication WO00/09552

Sequence 13323: Found in patent publication WO99/66041

AGTACTTCTAGAATTAATTAAGGCGGGGGAAAGTTAGAGGAACTGAAAGTTTGGGAATAG GCTGACCACATATTATGCCAGTGACCAGTATGACAGGAGATGGGGCCCTGCTGCCAGTCA 

Sequence 13324: Found in patent publication WO99/66041

ΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑ

Sequence 13325: Found in patent publication WO99/67383

AGTACTTCTAGAATTAATTAAGGCGGGTCCCTGCTTGTCCTATGAACTGCTCAGAGCTCC TGTCAGTCCAGCTGGGCCTTCTGGGTTCTGGCACCATTTCGTAGCCATTCTCTTTGTATT TTAAAAGGACGTTATGAAAGGGCTTAGACCAAAATAAATCATAATGTTACTTGAGCCACC TTATATAGCTGCTTGGAGAGTCTGTGTAGTTCTTTCTGCATGTATTAAAAATGTTTAGAA

Sequence 13326: Found in patent publication WO00/06698 GCCCTGGGAAACGATTGTGAACGCNTGAATGAATTGATGACTAANATCCGCTGCGGGGGT CCTACAGCGCANATGGTAATGCCCGTTCTGACTGGCTGGGAACGGCACCTTAGCAAGATA CTTAAAAGGCGCCTTCTGTGTGCCACNTGCACTCCACCCTGGGCAACAGAGCAAANACCC

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#### Table 3-6

CATCTCAAAAATAAATATATATAAAAAAATAAAAAGCTATTTCTAGTTTNATTTCAC TATAAAGTTTTGCTTTTATTAAA Sequence 13327: Found in patent publication US6034229 AGTACTTCTAGAATTAATTAAGGCGGGCGGATCATGAGGTCAGGAGTTCGAGACCAGCCT GGCCAACATGGTGAAACTCCGTGTCTACTAAAAATACAAAAATTGGCTGGGCCGTGGTGG AGGGTAGAGGTTGCAGTGAGCCAAGATTGTGCCACTGCACTCCAGCCTGGGCGACTGAGC AAGACTGTCTAAAAAAAAAAAAAAAAAATAAATAAAGNAA Sequence 13328: Found in patent publication WO00/18925 GCCGTCGTTTTACACCCCGNAANAANCNATANCNATGATNGNTGCCNGTCCCNCTCNNNA NAGNAATAATAAAANTAATTAANGCGGGTGATTAAATTTCAGTCCATTATGATTTTCCTT TCTCACATAATTACTTTTTCTTTTTAGACTTATAAGCTAGCAATTACAGATTTAACTAC AGCTATCAGCATGGACAAAAATAGTTATACAGCATTTTATAACAGAGCATTATGTTACAC CAAGATAAGGGAACTTCAAATGGCATTAACAGATTATGGAATTGTGCTGCTTCTTGATGC TACAGAAACTGTNAAACTAAATACCTTCCTTAATCGTGGACTCATCTACGTAGAACTAGG CCAGTATGGCTTTGCACTAGAGGATTTTAAAAC Sequence 15908: Found in patent publication WO98/33916 AAATTNNNTATTTATTTTAAANTAAAAANTATACTNAACTAATTATTTNAATTAAAGAAA Sequence 15909: Found in patent publication WO99/14328 Sequence 15910: Found in patent publication WO99/66041 Sequence 15911: Found in patent publication WO00/05382 AAGACATTCCAGACAAAAAGACTAACTTGTCAGAAAGCCCTGTGGCGGAAGGGAGCTTTT TTTGTAAAGGAGTTTTGGTTTTCTCCTAATAGCAATGGGATATCTTCCAAGGAATCTCAA TCAAAAGGGAGAGATGGCTCCGATTGGAATGTCATCCCTGGCTGAAGAGTAGAGGAAGCA AAAAAAAAAAAAAAAAAAAAAAANNGNNNNNNCCCNNTNTTTTTTT

## Table 4-1

Sequence ID	Accession	Sequence 52	AA321222
Sequence 1	AA371205	Sequence 53	AA482285
Sequence 2	Al366380	Sequence 54	AA933627
Sequence 3	AA315950	Sequence 55	AA349223
Sequence 4	AA112869	Sequence 56	AJ267257
Sequence 5	AA838003	Sequence 57	AA449435
Sequence 6	N52554	Sequence 58	AA029881
Sequence 7	AL039430	Sequence 59	Al267285
Sequence 8	A1620796	Sequence 60	AA449305
Sequence 9	Ai129714	Sequence 61	AA043133
Sequence 10	AA885293	Sequence 62	AL120521
Sequence 11	AA631903	Sequence 63	AA443154
Sequence 12	AI057093	Sequence 64	Al267185
Sequence 13	AA680351	Sequence 65	AA431868
Sequence 14	AA149579	Sequence 66	R28603
Sequence 15	H64252	Sequence 67	AI767050
Sequence 16	AA393816	Sequence 68	Al253345
Sequence 17	A1924963	Sequence 69	T08429
Sequence 18	R18412	Sequence 70	D53496
Sequence 19	AI267519	Sequence 71	Al267492
Sequence 20	AW068579	Sequence 72	AA868512
Sequence 21	AI051210	Sequence 73	T61421
Sequence 22	AA628113	Sequence 74	Al636911
Sequence 23	AI597980	Sequence 75	AI674594
Sequence 24	AA195579	Sequence 76	AA806717
Sequence 25	AA166952	Sequence 77	AA436125
Sequence 26	AA452288	Sequence 78	H17871
Sequence 27	AA188228	Sequence 79	AA075189
Sequence 28	AA256591	Sequence 80	AA147659
Sequence 29	R70639	Sequence 81	AA916753
Sequence 30	AA315689	Sequence 82	AA173990
Sequence 31	AA133268	Sequence 83	AI201294
Sequence 32	AA193411	Sequence 84	AA278445
Sequence 33	AA156244	Sequence 85	AI536588
Sequence 34	Al298496	Sequence 86	AA458549
Sequence 35	Al201298	Sequence 87	AA772075
Sequence 36	W07378	Sequence 88	A1499516
Sequence 37	AI089431	Sequence 89	AA156987
Sequence 38	AA649107	Sequence 90	AA493512
Sequence 39	AA714567	Sequence 91	AA522492
Sequence 40	A1744810	Sequence 92	AA331876
Sequence 41	AA024963	Sequence 93	R19183
Sequence 42	R14619	Sequence 94	AA381480
Sequence 43	AW104561	Sequence 95	AA625228
Sequence 44	AA296298	Sequence 96	A1793283
Sequence 45	AI267502	Sequence 97	Al694211
Sequence 46	AA804659	Sequence 98	AW023714
Sequence 47	AA053932	Sequence 99	AA004794
Sequence 48	A1366549	Sequence 100	AA027037
Sequence 49	Al267256	Sequence 101	R35797
Sequence 50	AA453996	Sequence 102	N31529
Sequence 51	AA482403	Sequence 103	AA578265

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Sequence 104	Al267255	Sequence 156	AA393650
Sequence 105	AI800250	Sequence 157	AA558281
Sequence 106	AA551189	Sequence 158	Z17864
Sequence 107	AA65 <b>2667</b>	Sequence 159	AL038162
Sequence 108	AA081218	Sequence 160	AA136944
Sequence 109	W21047	Sequence 161	AA493331
Sequence 110	Al216972	Sequence 162	N32192
Sequence 111	AA010986	Sequence 163	AA621405
Sequence 112	N72458	Sequence 164	AA485102
Sequence 113	AI089923	Sequence 165	AA262588
Sequence 114	AI039200	Sequence 166	AA760894
Sequence 115	AL121420	Sequence 167	Al692513
Sequence 116	Al123347	Sequence 168	AA312968
Sequence 117	AA654650	Sequence 169	AA305627
Sequence 118	Al417562	Sequence 170	AA336621
Sequence 119	AA047587	Sequence 171	AA521385
Sequence 120	AI954167	Sequence 172	AA055186
Sequence 121	AA167719	Sequence 173	R11877
Sequence 122	AA354341	Sequence 174	AA143021
Sequence 123	AA304846	Sequence 175	A1797353
Sequence 124	AA489 <b>44</b> 7	Sequence 176	AA234578
Sequence 125	AA634799	Sequence 177	D44737
Sequence 126	AI619798	Sequence 178	AA456721
Sequence 127	AI688807	Sequence 179	AA310142
Sequence 128	Al361119	Sequence 180	AA872858
Sequence 129	AA286965	Sequence 181	AA133742
Sequence 130	W37343	Sequence 182	AA028887
Sequence 131	A1188563	Sequence 183	AW022072
Sequence 132	W80673	Sequence 184	AA 152276
Sequence 133	AA564067	Sequence 185	AA443829
Sequence 134	AI033912	Sequence 186	AJ095031
Sequence 135	AI300489	Sequence 187	AA446789
Sequence 136	A1267454	Sequence 188	AA307857
Sequence 137	AA555115	Sequence 189	R10011
Sequence 138	AA524064	Sequence 190	AI267271
Sequence 139	AA808211	Sequence 191	AW148750
Sequence 140	AA248319	Sequence 192	A1692940
Sequence 141	AA679304	Sequence 193	AA469199
Sequence 142	AA018249	Sequence 194	AA070335
Sequence 143	AA303924	Sequence 195	AI873501
Sequence 144	AA311860	Sequence 196	AI769975
Sequence 145	AI624709	Sequence 197	AA522549
Sequence 146	AI219803	Sequence 198	AA142865
Sequence 147	AA149415	Sequence 199	AA431677
Sequence 148	AI630228	Sequence 200	AA349232
Sequence 149	AL047850	Sequence 201	AA521351
Sequence 150 Sequence 151	AA766817	Sequence 202	AI051959
•	A1933 <b>533</b>	Sequence 203	AA262123
Sequence 152	AA603158	Sequence 204	A1266043
Sequence 153	AA058906	Sequence 205	AA215332
Sequence 154	AA972598	Sequence 206	AA447189
Sequence 155	AI760 <b>699</b>	Sequence 207	AA037024

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Sequence 208	AW006468	Sequence 260	R49052
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Sequence 210	AA094656	Sequence 262	AI343404
Sequence 211	N40952	Sequence 263	AA625538
Sequence 212	AA165090	Sequence 264	AA134126
Sequence 213	AA158274	Sequence 265	AI188081
Sequence 214	AA196978	Sequence 266	AI015245
Sequence 215	AI631703	Sequence 267	AA460816
Sequence 216	AA679434	Sequence 268	A1557225
Sequence 217	AA306787	Sequence 269	AA156790
Sequence 218	AA708958	Sequence 270	AA397 <del>44</del> 8
Sequence 219	AA640474	Sequence 271	AA280952
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Sequence 221	R33762	Sequence 273	AA652600
Sequence 222	AA857478	Sequence 274	AA099572
Sequence 223	A1570611	Sequence 275	A1267384
Sequence 224	AA188312	Sequence 276	A1884829
Sequence 225	AA765909	Sequence 277	AA621165
Sequence 226	AA251627	Sequence 278	A1761728
Sequence 227	AA082063	Sequence 279	AA122237
Sequence 228	AI557626	Sequence 280	H51976
Sequence 229	AA204650	Sequence 281	Al391501
Sequence 230	AA335273	Sequence 282	AA613772
Sequence 231	Al346381	Sequence 283	AA5789 <b>7</b> 6
Sequence 232	D82303	Sequence 284	AA731507
Sequence 233	AI697849	Sequence 285	A1216358
Sequence 234	AA010869	Sequence 286	AI557112
Sequence 235	AA833543	Sequence 287	AL079529
Sequence 236	AA664472	Sequence 288	AA244003
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Sequence 238	N40262	Sequence 290	T59687
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Sequence 240	AI183745	Sequence 292	Al401220
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Sequence 242	AA587806	Sequence 294	AL047535
Sequence 243	AA355956	Sequence 295	AI267522
Sequence 244	AA225115	Sequence 296	A1682937
Sequence 245	AA504560	Sequence 297	F08579
Sequence 246	AA639999	Sequence 298	AI557053
Sequence 247	AA420680	Sequence 299	R83716
Sequence 248	R63251	Sequence 300	A1366467
Sequence 249	T67823	Sequence 301	H79449
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Sequence 251	R81053	Sequence 303	T06984
Sequence 252	AA457365	Sequence 304	H83781
Sequence 253	AA640280	Sequence 305	AA121145
Sequence 254	AA373628	Sequence 306	AA314941
Sequence 255	AA056744	Sequence 307	Al37959 <b>7</b>
Sequence 256	AL120828	Sequence 308	Al694477
Sequence 257	Al654123	Sequence 309	AA210833
Sequence 258	AA425414	Sequence 310	AA521163
Sequence 259	D60110	Sequence 311	AA126834
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Sequence 312	AA437224	Sequence 364	AA659719
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Sequence 314	AA436197	Sequence 366	AA235224
Sequence 315	AA313692	Sequence 367	AI018034
Sequence 316	AA836682	Sequence 368	AA284997
Sequence 317	AA029276	Sequence 369	AA401802
Sequence 318	AA017175	Sequence 370	T71615
Sequence 319	AA516531	Sequence 371	R23858
Sequence 320	AA465494	Sequence 372	AL039698
Sequence 321	Al394646	Sequence 373	AF017714
Sequence 322	AA927118	Sequence 374	A1267570
Sequence 323	AW062373	Sequence 375	AA525386
Sequence 324	Z20613	Sequence 376	AL037256
Sequence 325	AA134311	Sequence 377	A1040598
Sequence 326	AA612864	Sequence 378	W58731
Sequence 327	AA631868	Sequence 379	A1356805
Sequence 328	Al239435	Sequence 380	AA033951
Sequence 329	T70288	Sequence 381	AA282215
Sequence 330	AA285043	Sequence 382	AA421795
Sequence 331	AA705546	Sequence 383	AA767784
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Sequence 333	AI082617	Sequence 385	R18215
Sequence 334	AA345027	Seque <b>nce 386</b>	AA122080
Sequence 335	Al217172	Sequence 387	AA157875
Sequence 336	AA449907	Sequence 388	AA040673
Sequence 337	AA626511	Sequence 389	AA463761
Sequence 338	Al002821	Sequence 390	AA527805
Sequence 339	AA505073	Sequence 391	AA036798
Sequence 340	T71154	Sequence 392	R25807
Sequence 341	AW070326	Sequence 393	AA675892
Sequence 342	AA366269	Sequence 394	W03191
Sequence 343	R71619	Sequence 395	AI016031
Sequence 344	AA058568	Sequence 396	AA258079
Sequence 345	AA359418	Sequence 397	AA150033
Sequence 346	AA305599	Sequence 398	AA297647
Sequence 347	Al269053	Sequence 399	Al610699
Sequence 348	AA403121	Sequence 400	AA120964
Sequence 349	AA514302	Sequence 401	AA814091
Sequence 350	AA174144	Sequence 402	Al474235
Sequence 351	AW177204	Sequence 403	AA307371
Sequence 352	AA580097	Sequence 404	AA344133
Sequence 353	AA664480	Sequence 405	AA316649
Sequence 354	W60040	Sequence 406	A1694320
Sequence 355	R71431	Sequence 407	AA017096
Sequence 356	AA581846	Sequence 408	AA452999
Sequence 357	Al216968	Sequence 409	AA297472
Sequence 358	AA243573	Sequence 410	W87509
Sequence 359	AL037638	Sequence 411	AA020941
Sequence 360	R497 <b>34</b>	Sequence 412	R64194
Sequence 361	AA223433	Sequence 413	A1042293
Sequence 362	AA157267	Sequence 414	AA044118
Sequence 363	AL046802	Sequence 415	H25806

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Sequence 416	AA383011	Sequence 468	AA308486
Sequence 417	AA328366	Sequence 469	AA773998
Sequence 418	Al216969	Sequence 470	AA644237
Sequence 419	AA165339	Sequence 471	AA173721
Sequence 420	AA308063	Sequence 472	AA458528
Sequence 421	N52271	Sequence 473	AI400752
Sequence 422	AA315718	Sequence 474	AA807276
Sequence 423	A1554809	Sequence 475	AW022792
Sequence 424	H81755	Sequence 476	A1989600
Sequence 425	AA829350	Sequence 477	Al267838
Sequence 426	R45470	Sequence 478	AA825265
Sequence 427	AA133652	Sequence 479	AA229199
Sequence 428	AA405892	Sequence 480	AW003442
Sequence 429	AW139567	Sequence 481	AI275711
Sequence 430	AA225114	Sequence 482	R11581
Sequence 431	AA343168	Sequence 483	AA931943
Sequence 432	AA040405	Sequence 484	AA7702 <b>72</b>
Sequence 433	AA229745	Sequence 485	D55192
Sequence 434	AA830959	Sequence 486	AA129551
Sequence 435	AW075906	Sequence 487	AA772683
Sequence 436	AA533227	Sequence 488	AA047752
Sequence 437	AA079041	Sequence 489	Al366471
Sequence 438	N31825	Sequence 490	Al267 <b>66</b> 9
Sequence 439	AA503132	Sequence 491	AA702679
Sequence 440	AA318969	Sequence 492	AA449544
Sequence 441	AW020429	Sequence 493	AA133757
Sequence 442	AA315072	Sequence 494	AA287347
Sequence 443	AA709208	Sequence 495	AA548129
Sequence 444	H14446	Sequence 496	AA355351
Sequence 445	AA548124	Sequence 497	AA164703
Sequence 446	AI927636	Sequence 498	H15660
Sequence 447	Al697249	Sequence 499	A1417910
Sequence 448	AA146900	Sequence 500	AL041265
Sequence 449	N94259	Sequence 501	A1281449
Sequence 450	AA215284	Sequence 502	AA447021
Sequence 451	A1000877	Sequence 503	AA032194
Sequence 452	AA373289	Sequence 504	AI079595
Sequence 453	AA040624	Sequence 505	AA181179
Sequence 454	AA227789	Sequence 506	AA368792
Sequence 455	Al378932	Sequence 507	AA5228 <b>5</b> 6
Sequence 456	AA936532	Sequence 508	A1004135
Sequence 457	AL043113	Sequence 509	D82197
Sequence 458	AL121031	Sequence 510	AW083338
Sequence 459	AA508810	Sequence 511	AI076608
Sequence 460	AA129003	Sequence 512	A1754013
Sequence 461	R63823	Sequence 513	AA088322
Sequence 462	N24785	Sequence 514	A1480424
Sequence 463	T95139	Sequence 515	AA151796
Sequence 464	A1480345	Sequence 516	AA034322
Sequence 465	AA181111	Sequence 517	A1207660
Sequence 466	AA908961	Sequence 518	T81369
Sequence 467	AA836233	Sequence 519	AA885902

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Sequence 520	A1332597	Sequence 572	AI198986
Sequence 521	AA043891	Sequence 573	A1972279
Sequence 522	AA531606	Sequence 574	H00112
Sequence 523	AA504728	Sequence 575	AI267480
Sequence 524	AA315118	Sequence 576	A1623984
Sequence 525	Al391717	Sequence 577	AA701667
Sequence 526	AA337114	Sequence 578	AA988520
Sequence 527	AA573557	Sequence 579	AA312025
Sequence 528	A1630584	Sequence 580	AA421510
Sequence 529	AA534235	Sequence 581	AA259166
Sequence 530	H56898	Sequence 582	AA459831
Sequence 531	AI073907	Sequence 583	AA284410
Sequence 532	Al471561	Sequence 584	AA578295
Sequence 533	T28974	Sequence 585	R20651
Sequence 534	AA307475	Sequence 586	AA364882
Sequence 535	A1952763	Sequence 587	AA135135
Sequence 538	AA524893	Sequence 588	AI332939
Sequence 537	Al927846	Sequence 589	W87865
Sequence 538	H12056	Sequence 590	AA465598
Sequence 539	AA010758	Sequence 591	AA022478
Sequence 540	AA334403	Sequence 592	U72041
Sequence 541	A1625404	Sequence 593	AI239822
Sequence 542	Al356309	Sequence 594	Al253293
Sequence 543	A1768908	Sequence 595	AA206721
Sequence 544	T06925	Sequence 596	AA447322
Sequence 545	AI656158	Sequence 597	AA044896
Sequence 546	AI366461	Sequence 598	AA149421
Sequence 547	T30677	Sequence 599	AL047056
Sequence 548	AA436472	Sequence 600	AW057515
Sequence 549	AA628409	Sequence 601	AA774247
Sequence 550	AA263162	Sequence 602	AA412501
Sequence 551	AA468167	Sequence 603	Al458457
Sequence 552	Al929464	Sequence 604	AA307513
Sequence 553	AA465340	Sequence 605	AI761702
Sequence 554	A1984636	Sequence 606	AA400418
Sequence 555	AA687571	Sequence 607	AA524651
Sequence 556	AA488809	Sequence 608	AW102941
Sequence 557	AI557235	Sequence 609	AW188391
Sequence 558	AA609641	Sequence 610	T07196
Sequence 559	Al458673	Sequence 611	AI440194
Sequence 560	AW166338	Sequence 612	A1431350
Sequence 561	AA984384	Sequence 613	R60395
Sequence 562	AI923011	Sequence 614	AA263042
Sequence 563	A1744818	Sequence 615	AI267216
Sequence 564	AA708501	Sequence 616	AA027868
Sequence 565	AA402500	Sequence 617	AI357642
Sequence 566	AA132633	Sequence 618	AA430752
Sequence 567	AI523222	Sequence 619	AI827667
Sequence 568	AA814859	Sequence 620	AA135254
Sequence 569	H54240	Sequence 621	AW067913
Sequence 570	Al206223	Sequence 622	AI914128
Sequence 571	Al627818	Sequence 623	AA430313
ocquerios o/ I	7.1027010	Sequence 023	, VA-1000 13

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			A106000E
Sequence 624	AA047517	Sequence 676	Al253335 AL045332
Sequence 625	AA075810	Sequence 677	AA228953
Sequence 626	AA001414	Sequence 678	AA535837
Sequence 627	AA375531	Sequence 679	AW020820
Sequence 628	R51416	Sequence 680	A1308072
Sequence 629	AA244066	Sequence 681	AA975275
Sequence 630	AA550927	Sequence 682	AI041116
Sequence 631	AA507791	Sequence 683	AI433757
Sequence 632	AI002745	Sequence 684	AI753951
Sequence 633	AA285290	Sequence 685	AA809784
Sequence 634	AI253379	Sequence 686	AA459476
Sequence 635	AA024480	Sequence 687	
Sequence 636	AA595047	Sequence 688	W37862
Sequence 637	AA281307	Sequence 689	AA206147
Sequence 638	AA468760	Sequence 690	A1697873
Sequence 639	AI670913	Sequence 691	AA311772
Sequence 640	AA311639	Sequence 692	AA447815
Sequence 641	Al128986	Sequence 693	AA312591
Sequence 642	AA478891	Sequence 694	AA216433
Sequence 643	AA613935	Sequence 695	AA622418
Sequence 644	AA318009	Sequence 696	AI922040
Sequence 645	AA203537	Sequence 697	AI589206
Sequence 646	AA017262	Sequence 698	AI928622
Sequence 647	AA351443	Sequence 699	AA485767
Sequence 648	AA352938	Sequence 700	AA070393
Sequence 649	AA446322	Sequence 701	Al216966
Sequence 650	F12000	Sequence 702	AA639964
Sequence 651	AA001460	Sequence 703	AA204847
Sequence 652	AA346504	Sequence 704	AA491866
Sequence 653	AA779728	Sequence 705	AA227647
Sequence 654	AA349541	Sequence 706	AA448514
Sequence 655	AA115636	Sequence 707	AA171807
Sequence 656	A1050871	Sequence 708	AA765140
Sequence 657	AA827196	Sequence 709	AA172188
Sequence 658	AA393886	Sequence 710	AA029835 AA907660
Sequence 659	AA083288	Sequence 711	AA770505
Sequence 660	AA302548	Sequence 712	AA070624
Sequence 661	AI085559	Sequence 713	AI139200
Sequence 662	AA812209	Sequence 714	AA025834
Sequence 663	A1253347	Sequence 715	AA504322
Sequence 664	AA082629	Sequence 716	AA307033
Sequence 665	AA121656	Sequence 717	Al269415
Sequence 666	R88370	Sequence 718	
Sequence 667	AA383100	Sequence 719	AA345139
Sequence 668		Sequence 720	AA361618
Sequence 669	AA280873	Sequence 721	AA627073
Sequence 670		Sequence 722	R74152 AA578773
Sequence 671		Sequence 723	
Sequence 672		Sequence 724	AA558636
Sequence 673		Sequence 725	
Sequence 674	Z44402	Sequence 726	
Sequence 675	Al366374	Sequence 727	H12519

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Sequence 728	AA508889	Sequence 780	AA224269
Sequence 729	AL036423	Sequence 781	A:708364
Sequence 730	AA135065	Sequence 782	AA442352
Sequence 731	W93718	Sequence 783	AA725525
Sequence 732	AA022788	Sequence 784	AA418403
Sequence 733	AA358369	Sequence 785	AA228845
Sequence 734	F12883	Sequence 786	AA453 <b>550</b>
Sequence 735	AA523902	Sequence 787	AA522664
Sequence 736	AA255526	Sequence 788	AI133612
Sequence 737	W94398	Sequence 789	AW138413
Sequence 738	AA435877	Sequence 790	Al216986
Sequence 739	T83709	Sequence 791	A:638492
Sequence 740	R14684	Sequence 792	AW103377
Sequence 741	Al871555	Sequence 793	AA486413
Sequence 742	AA037683	Sequence 794	AA019152
Sequence 743	Al188078	Sequence 795	H64556
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Sequence 745	AA364656	Sequence 797	AA370976
Sequence 746	AA489433	Sequence 798	AA502798
Sequence 747	AA037450	Sequence 799	AA446366
Sequence 748	AA460930	Sequence 800	N67168
Sequence 749	AA166925	Sequence 801	W04240
Sequence 750	N98749	Sequence 802	AA126750
Sequence 751	AA417324	Sequence 803	H67975
Sequence 752	AA620750	Sequence 804	Al005449
Sequence 753	AA744849	Sequence 805	AA947511
Sequence 754	AA507041	Sequence 806	AL120396
Sequence 755	AI040234	Sequence 807	AA229495
Sequence 756	AA228940	Sequence 808	AA324252
Sequence 757	F05508	Sequence 809	AA354245
Sequence 758	C04925	Sequence 810	AA314028
Sequence 759	AA662702	Sequence 811	AA574248
Sequence 760	Al630144	Sequence 812	AA569482
Sequence 761	T72376	Sequence 813	AA226321
Sequence 782	AA225438	Sequence 814	AI025519
Sequence 763	AA190674	Sequence 815	AA724392
Sequence 764	AA070031	Sequence 816	AA125809
Sequence 765	AA526368	Sequence 817	AA843176
Sequence 766	AA357593	Sequence 818	AW023001
Sequence 767	AA527730	Sequence 819	Al338190
Sequence 768	W26780	Sequence 820	AI817242
Sequence 769	AA002131	Sequence 821	AI521964
Sequence 770	Al332588	Sequence 822	AA325580
Sequence 771	R92548	Sequence 823	AA347577
Sequence 772	AA229713	Sequence 824	D56420
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Sequence 775	AA188396	Sequence 827	W27526
Sequence 776	AA452412	Sequence 828	Al380812
Sequence 777	AA112710	Sequence 829	AA625145
Sequence 778	AA385204	Sequence 830	AA573974
Sequence 779	AA295737	Sequence 831	AI525651
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Sequence 832	AI791719		R14689
Sequence 833	N40717	Sequence 885	AA364190
Sequence 834	Al694767	Sequence 886	AA044750
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Sequence 838	AA633439	Sequence 890	AA314920
Sequence 839	AA847597	Sequence 891	AA689334
Sequence 840	Al312562	Sequence 892	
Sequence 841	AA931666	Sequence 893	W03366 AA405068
Sequence 842	AA742701	Sequence 894	AA417904
Sequence 843	R69818	Sequence 895	
Sequence 844	AI379679	Sequence 896	AA347085
Sequence 845	W19759	Sequence 897	AA702738
Sequence 846	AA683080	Sequence 898	AA088606
Sequence 847	AA972883	Sequence 899	AA528202
Sequence 848	AA722090	Sequence 900	C75042
Sequence 849	Al267490	Sequence 901	H92683
Sequence 850	Al110836	Sequence 902	AA602675
Sequence 851	AI064763	Sequence 903	AA146941
Sequence 852	AA627785	Sequence 904	AA923172
Sequence 853	AL036650	Sequence 905	A1470259
Sequence 854	AA278473	Sequence 906	AA722204
Sequence 855	AA515109	Sequence 907	F06621
Sequence 856	AL119859	Seque <b>nce 908</b>	AA090790
Sequence 857	AA280221	Sequen <b>ce 909</b>	H05326
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Sequence 859	AW178543	Sequence 911	AA298549
Sequence 860	A1826504	Sequence 912	AL036170
Sequence 861	A1754874	Sequence 913	T06574
Sequence 862	A1948822	Sequence 914	AA298285
Sequence 863	AA188875	Sequence 915	AA326242
Sequence 864	AA988011	Sequence 916	C20630
Sequence 865	AA489965	Sequence 917	AA226266
Sequence 866	AA946679	Sequence 918	AA486041
Sequence 867	AA522850	Sequence 919	AA225154
Sequence 868	AI813724	Sequence 920	F09972
Sequence 869	A1740796	Sequence 921	AI590351
Sequence 870	AA317662	Sequence 922	AA301408
Sequence 871	AA099878	Sequence 923	AA505423
Sequence 872	AA372911	Sequence 924	AI693302
Sequence 873	T68793	Sequence 925	H09642
Sequence 874	AA469129	Sequence 926	AA146589
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		Sequence 928	AA044599
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Sequence 878		Sequence 931	AA057867
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Sequence 880		Sequence 933	AA112441
Sequence 881		Sequence 934	AA040135
Sequence 882		Sequence 935	AA480124
Sequence 883	AAU/5012	ocquenice 000	

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Sequence 936	AA233943	Sequence 988	AI880872
Sequence 937	AA306902	Sequence 989	T30767
Sequence 938	AA319070	Sequence 990	AI174822
Sequence 939	AA083370	Sequence 991	AI674871
Sequence 940	AI458823	Sequence 992	AA838697
Sequence 941	Al168205	Sequence 993	AA432264
Sequence 942	AA935798	Sequence 994	AI199681
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Sequence 944	AA069880	Sequence 996	N67109
Sequence 945	AA707511	Sequence 997	AA169493
Sequence 946	Al263110	Sequence 998	AI554590
Sequence 947	AA713530	Sequence 999	R59789
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Sequence 949	AA576669	Sequence 1001	AW117546
Sequence 950	AA664798	Sequence 1002	AA029403
Sequence 951	A1696488	Sequence 1003	AI248187
Sequence 952	AI758378	Sequence 1004	A1370846
Sequence 953	AJ741143	Sequence 1005	AA069801
Sequence 954	AA430271	Sequence 1006	AL120694
Sequence 955	AJ754461	Sequence 1007	H83524
Sequence 956	AI280935	Sequence 1008	AI948503
Sequence 957	AA541813	Sequence 1009	AA631437
Sequence 958	AA377796	Sequence 1010	A1744509
Sequence 959	C02739	Sequence 1011	A1050966
Sequence 960	AA165562	Sequence 1012	AA418177
Sequence 961	AL036801	Sequence 1013	AA644624
Sequence 962	AA045870	Sequence 1014	AA025551
Sequence 963	AA300869	Sequence 1015	R42583
Sequence 964	AA305030	Sequence 1016	AI357346
Sequence 965	AA761069	Sequence 1017	AL040518
Sequence 966	AA641623	Sequence 1018	AI627356
Sequence 967	Al246615	Sequence 1019	AA056058
Sequence 968	AA299140	Sequence 1020	AA993414
Sequence 969	AA847452	Sequence 1021	AW129657
Sequence 970	R50277	Sequence 1022	AA399265
Sequence 971	AA252436	Sequence 1023	AA102274
Sequence 972	AA016116	Sequence 1024	N28426
Sequence 973	AA909045	Sequence 1025	AA069407
Sequence 974	AA713517	Sequence 1026	AA652478
Sequence 975	AA743453	Sequence 1027	AA126951
Sequence 976	AL048852	Sequence 1028	AA813266
Sequence 977	AI298459	Sequence 1029	Al309121
Sequence 978	N40598	Sequence 1030	AA743710
Sequence 979	AW173161	Sequence 1031	AA229424
Sequence 980	AA378508	Sequence 1032	AA334754
Sequence 981	AA548465	Sequence 1033	AA401469
Sequence 982	AA777529	Sequence 1034	AL046756
Sequence 983	AA176979	Sequence 1035	AA680052
Sequence 984	AA506459	Sequence 1036	AA632291
Sequence 985	AA306232	Sequence 1037	AA349192
Sequence 986	AA280917	Sequence 1038	AI948513
Sequence 987	Al690979	Sequence 1039	AA772465

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Sequence 1040	AA129772	00400	1143623
Sequence 1041	AI270536	Ccqucilor loss	W169473
Sequence 1042	Al124856	204251102 1001	A205647
Sequence 1043	AA244071	204201100 1000	1755085
Sequence 1044	Al253319	· · · · · · · · · · · · · · ·	1627945
Sequence 1045	Al188850	Ocquerios res	A599864
Sequence 1046	AA083300	Codasiiss	A403109
Sequence 1047	AA977226	00400.00	A4767 <b>56</b>
Sequence 1048	AA121615	Ocquerios (1700)	A532883
Sequence 1049	Al955087	Sequence	A305697
Sequence 1050	Al332459	Ccquonise	N355050
Sequence 1051	AA506299	ocquomos	006419
Sequence 1052	Al434180	004001100 1101	AA247232
Sequence 1053	AA102698	Sequence 1105	41092766
Sequence 1054	AA594131	Ocquemes	AA705269
Sequence 1055	AA806057	— — — — — — — — — — — — — — — — — — —	AA34592 <b>5</b>
Sequence 1056	Al683431	Sequence 1108	AA010828
Sequence 1057	AA181017	Sequence 1109	Al302083
Sequence 1058	W37322	Sequence 1110	AA011000
Sequence 1059	AI949242	Sequence 1111	A1080480
Sequence 1060	AA148087		AA984407
Sequence 1061	AA053917	Sequence 1113	AA405758
Sequence 1062	AA837048	Sequence 1114	Al373634
Sequence 1063	AI743954	Sequence 1115	Al459234
Sequence 1064	Al623891	Sequence 1116	AA481476
		Sequence 1117	Al278611
Sequence 1065 Sequence 1066		Sequence 1118	M78906
Sequence 1067		Sequence 1119	A1031653
Sequence 1068		Sequence 1120	AF150208
Sequence 1069		Sequence 1121	AA525487
Sequence 1070		Sequence 1122	AA351741
Sequence 1071		Sequence 1123	AW073731
Sequence 1072		Sequence 1124	AA430160
Sequence 1073		Sequence 1125	A1084005
Sequence 1074		Sequence 1126	AA157733
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Sequence 107		Sequence 1131	AA314161
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Sequence 108		Sequence 1135	A1022340
Sequence 108		Sequence 1136	A1802946
Sequence 108		Sequence 1137	AA479284
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Sequence 100		Sequence 1139	
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Sequence 10		Sequence 1142	A1869322
Sequence 10		Sequence 1143	AA773478
Sequence to	<del></del>		

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Sequence 1144	AA968556	Sequence 1196	Z25169
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Sequence 1146	AA608749	Sequence 1198	AA022937
Sequence 1147	AA521213	Sequence 1199	AA255974
Sequence 1148	AA430754	Sequence 1200	AA625467
Sequence 1149	AA917666	Sequence 1201	AA602662
Sequence 1150	AI743852	Sequence 1202	Al420026
Sequence 1151	AA004642	Sequence 1203	AA215538
Sequence 1152	AA531563	Sequence 1204	AI133208
Sequence 1153	AA280753	Sequence 1205	R11114
Sequence 1154	AA9068 <b>68</b>	Sequence 1206	AA479689
Sequence 1155	AA479781	Sequence 1207	AA528104
Sequence 1158	AA446116	Sequence 1208	AA034912
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Sequence 1160	Al267321	Sequence 1212	AA307209
Sequence 1161	AA679068	Sequence 1213	Al332712
Sequence 1162	AI557336	Sequence 1214	AA702942
Sequence 1163	AA460440	Sequence 1215	AW163314
Sequence 1164	AA308091	Sequence 1216	AA336412
Sequence 1165	AA513597	Sequence 1217	H58571
Sequence 1166	AA442287	Sequence 1218	AA186364
Sequence 1167	AW084125	Sequence 1219	AA398463
Sequence 1168	Al274700	Sequence 1220	AA017500
Sequence 1169	AW138181	Sequence 1221	AW088608
Sequence 1170	AA576425	Sequence 1222	AA493183
Sequence 1171	AA304651	Sequence 1223	Al240899
Sequence 1172	A1285435	Sequence 1224	R55600
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Sequence 1176	AI340044	Sequence 1228	AA188771
Sequence 1177	AL039293	Sequence 1229	AA234050
Sequence 1178	C18181	Sequence 1230	Al557495
Sequence 1179	AA449649	Sequence 1231	AW080196
Sequence 1180	AA312605	Sequence 1232	AA418719
Sequence 1181	AA046577	Sequence 1233	AA225294
Sequence 1182	A1075037	Sequence 1234	AA812993
Sequence 1183	Al421027	Sequence 1235	AA312689
Sequence 1184	AA781086	Sequence 1236	H13891
Sequence 1185	AA825512	Sequence 1237	T77631
Sequence 1186	AA205546	Sequence 1238	AA311905
Sequence 1187	AA027882	Sequence 1239	AI203546
Sequence 1188	AA810418	Sequence 1240	AJ435606
Sequence 1189	AA348878	Sequence 1241	AA143170
Sequence 1190	AA281653	Sequence 1242	AL037316
Sequence 1191	H17170	Sequence 1243	AJ205868
Sequence 1192	AA195751	Sequence 1244	AW103458
Sequence 1193	AA364261	Sequence 1245	W90519
Sequence 1194	AA235838	Sequence 1246	Al367641
Sequence 1195	AA503943	Sequence 1247	AA394298

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Sequence 1248	AA167826	Sequence 1300	U14528
Sequence 1249	AA322540	Sequence 1301	AF153608
Sequence 1250	Al253192	Sequence 1302	U57847
Sequence 1251	AA843531	Sequence 1303	AL049276
Sequence 1252	AA203110	Sequence 1304	AF124438
Sequence 1253	Al690278	Sequence 1305	AF007144
Sequence 1254	AA506323	Sequence 1306	L01100
Sequence 1255	AA406 <b>478</b>	Sequence 1307	AF056617
Sequence 1256	Al816058	Sequence 1308	U57877
Sequence 1257	AA149219	Sequence 1309	X74331
Sequence 1258	AW176020	Sequence 1310	S82081
Sequence 1259	AA507792	Sequence 1311	AF115402
Sequence 1260	AA468839	Sequence 1312	M29366
Sequence 1261	AA195151	Sequence 1313	X04106
Sequence 1262	AA070277	Sequence 1314	AF151103
Sequence 1263	T08287	Sequence 1315	S <b>75755</b>
Sequence 1264	AA130555	Sequence 1316	X76057
Sequence 1265	AL122121	Sequence 1317	AF070641
Sequence 1266	AF112481	Sequence 1318	X78933
Sequence 1267	AF051334	Sequence 1319	AF003083
Sequence 1268	U61397	Sequence 1320	S <b>73498</b>
Sequence 1269	AF116827	Sequence 1321	AB022663
Sequence 1270	AB029020	Sequence 1322	X94910
Sequence 1271	AL110183	Sequence 1323	X01742
Sequence 1272	AB020636	Sequence 1324	AB020633
Sequence 1273	AF000986	Sequence 1325	D14043
Sequence 1274	D86326	Sequence 1326	L10284
Sequence 1275	AJ005016	Sequence 1327	X15729
Sequence 1276	AB020637	Sequence 1328	AB020680
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Sequence 1278	D21209	Sequence 1330	<b>⊔77396</b>
Sequence 1279	L070 <b>77</b>	Sequence 1331	U84138
Sequence 1280	X74801	Sequence 1332	AB007873
Sequence 1281	AB015333	Sequence 1333	D87442
Sequence 1282	AB011119	Sequence 1334	D42045
Sequence 1283	Y11395	Sequence 1335	AF008442
Sequence 1284	AB029028	Sequence 1336	AJ002955
Sequence 1285.	AF054179	Sequence 1337	M94856
Sequence 1286	M21895	Sequence 1338	X16940
Sequence 1287	AL080206	Sequence 1339	M36634
Sequence 1288	AL050355	Sequence 1340	AB018266
Sequence 1289	AB006965	Sequence 1341	M17323
Sequence 1290	AL117563	Sequence 1342	X75593
Sequence 1291	X75861	Sequence 1343	J04137
Sequence 1292	D13315	Sequence 1344	AL049969
Sequence 1293	D50420	Sequence 1345	D13641
Sequence 1294		Sequence 1346	D31886
Sequence 1295		Sequence 1347	X63563
Sequence 1296		Sequence 1348	D63480
Sequence 1297		Sequence 1349	AF039918
Sequence 1298		Sequence 1350	X80910
Sequence 1299		Sequence 1351	D87969
22422,100 ,200		<b>-</b>	

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Sequence 1352	Z24724	Sequence 1404	AB019435
Sequence 1353	U47674	Sequence 1405	L41143
Sequence 1354	AB011158	Sequence 1406	AF141327
Sequence 1355	AL050071	Sequence 1407	U80034
Sequence 1356	AB014610	Sequence 1408	AF062318
Sequence 1357	AL050290	Sequence 1409	Y13834
Sequence 1358	AD001528	Sequence 1410	AB024518
Sequence 1359	U67784	Sequence 1411	AB007898
Sequence 1360	L09749	Sequence 1412	U588 <b>55</b>
Sequence 1361	X85129	Sequence 1413	K03515
Sequence 1362	AF022913	Sequence 1414	M81757
Sequence 1363	AF131738	Sequence 1415	M60527
Sequence 1364	AJ004913	Sequence 1416	AL110181
Sequence 1365	A8002383	Sequence 1417	M25756
Sequence 1366	J03015	Sequence 1418	AB033061
Sequence 1367	D29954	Sequence 1419	D31770
Sequence 1368	AF069469	Sequence 1420	AB004066
Sequence 1369	AL049996	Sequence 1421	E01932
Sequence 1370	D42044	Sequence 1422	U90909
Sequence 1371	AL080181	Sequence 1423	AF078864
Sequence 1372	AF061258	Sequence 1424	AF047020
Sequence 1373	AL110153	Sequence 1425	J04183
Sequence 1374	AL080202	Sequence 1426	AL050089
Sequence 1375	U19769	Sequence 1427	X57025
Sequence 1376	AB003103	Sequence 1428	D14710
Sequence 1377	L13848	Sequence 1429	A18585
Sequence 1378	M24194	Sequence 1430	AL049932
Sequence 1379	J05 <b>480</b>	Sequence 1431	Z50749
Sequence 1380	M16342	Sequence 1432	AB014579
Sequence 1381	AB002365	Sequence 1433	AF000652
Sequence 1382	AF144566	Sequence 1434	X03559
Sequence 1383	AJ001443	Sequence 1435	L37080
Sequence 1384	U54559	Sequence 1436	M29548
Sequence 1385	AF075061	Sequence 1437	AB000468
Sequence 1386	U37143	Sequence 1438	AF061261
Sequence 1387	D79986	Sequence 1439	AF126780
Sequence 1388	AF013758	Sequence 1440	X77196
Sequence 1389	AB020694	Sequence 1441	A3007949
Sequence 1390	AF103804	Sequence 1442	D49490
Sequence 1391	AF052578	Sequence 1443	AF006086
Sequence 1392	AL117599	Sequence 1444	U39657
Sequence 1393	AL110211	Sequence 1445	AF005043
Sequence 1394	AB018342	Sequence 1446	AF053535
Sequence 1395	AF151844	Sequence 1447	AL110141
Sequence 1396	AL050192	Sequence 1448	AF151805
Sequence 1397	X97065	Sequence 1449	AB033049
Sequence 1398	L12168	Sequence 1450	L08437
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Sequence 1400	X59405	Sequence 1452	M74777
Sequence 1401	X76732	Sequence 1453	L31801
Sequence 1402	U36336	Sequence 1454	X60708
Sequence 1403	AF131802	Sequence 1455	AF052124

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Sequence 1456 AF077202	Sequence 1508 M74002
Sequence 1457 E01915	Sequence 1509 D87735
Sequence 1458 M22865	Sequence 1510 L17128
Sequence 1459 M73837	Sequence 1511 D21163
Sequence 1460 U72937	Sequence 1512 AF070523
Sequence 1461 S72481	Sequence 1513 AB011148
Sequence 1462 AB020724	Sequence 1514 L13385
Sequence 1463 AL117621	Sequence 1515 D13988
Sequence 1464 ABO19568	Sequence 1516 AL110238
Sequence 1465 L05425	Sequence 1517 D21210
Sequence 1466 AL122097	Sequence 1518 AB011087
Sequence 1467 AF151868	Sequence 1519 AF054284
Sequence 1468 AF085355	Sequence 1520 AF176702
Sequence 1469 AL049471	Sequence 1521 AB004857
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Sequence 1471 AB007956	Sequence 1523 AF035293
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Sequence 1475 U16850	Sequence 1527 AF001433
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Sequence 1477 AB026723	Sequence 1529 AF131775
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Sequence 1481 Y09305	Sequence 1533 AF038451
Sequence 1482 M14648	Sequence 1534 D00422
Sequence 1483 AF126181	Sequence 1535 D13388
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Sequence 1485 D87845	Sequence 1537 L16510
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Sequence 1487 D26485	Sequence 1539 E02628
Sequence 1488 X73608	Sequence 1540 U62434
Sequence 1489 U64820	Sequence 1541 J02923
Sequence 1490 D89053	Sequence 1542 U60975
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Sequence 1492 AJ002744	Sequence 1544 AF167572
Sequence 1493 D31883	Sequence 1545 AF169797
Sequence 1494 U01923	Sequence 10 to
Sequence 1495 AB018331	Sequence 1011
Sequence 1496 AB019563	Sequence 19 19
Sequence 1497 U41813	Sequence 10.10
Sequence 1498 M61199	Sequence 1000
Sequence 1499 M22538	Sequence 1001
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Sequence 1501 AL049944	
Sequence 1502 AB007963	Sequence 1001
Sequence 1503 AB006968	Seducios (ess
Sequence 1504 AF077037	Sequence 1202
Sequence 1505 U43701	Sequence 1007
Sequence 1506 E05957	0040000
Sequence 1507 AF103907	Sequence 1559 D50372

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Sequence 1560	D12676	Sequence 1612	U39064
Sequence 1561	AF098297	Sequence 1613	U17714
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Sequence 1564	U53468	Sequence 1616	AF100759
Sequence 1565	AF165281	Sequence 1617	AB019564
Sequence 1566	U12979	Sequence 1618	D63998
Sequence 1567	AF188746	Sequence 1619	AL050265
Sequence 1568	D49489	Sequence 1620	AB033078
Sequence 1569	M24630	Sequence 1621	X77548
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Sequence 1587	D50645	Sequence 1639	D63486
Sequence 1588	AF131749	Sequence 1640	AJ000644
Sequence 1589	AF013759	Sequence 1641	L38961
Sequence 1590	AF097535	Sequence 1642	AF042166
Sequence 1591	Y12860	Sequence 1643	AF078847
Sequence 1592	X60489	Sequence 1644	AJ130733
Sequence 1593	Z48042	Sequence 1645	X54326
Sequence 1594	U79751	Sequence 1646	J04208
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Sequence 1599	U03274	Sequence 1651	E03414
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Sequence 1602	J04794	Sequence 1654	M61831
Sequence 1603	AF039029	Sequence 1655	U39360
Sequence 1604	Z22534	Sequence 1656	AL050255
Sequence 1605	M10036	Sequence 1657	U73824
Sequence 1606	AF131831	Sequence 1658	Y00281
Sequence 1607	AF068754	Sequence 1659	U41515
Sequence 1608	AL049367	Sequence 1660	AF100761
Sequence 1609 Sequence 1610	X53793	Sequence 1661	AF151855
Sequence 1611	AF070553	Sequence 1662	AB007618
ocdine (ne (o))	AB018281	Sequence 1663	AF055994

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Sequence 1664	AF047472	Sequence 1716	Y12065
Sequence 1665		Sequence 1717	M85168
Sequence 1666	AJ237946	Sequence 1718	L11066
Sequence 1667	E08293	Sequence 1719	AF016270
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Sequence 1672	AF070561	Sequence 1724	U68536
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Sequence 1674	M74524	Sequence 1726	D87667
Sequence 1675	AB000220	Sequence 1727	AB028973
Sequence 1676	AB020639	Sequence 1728	AF106943
Sequence 1677	AF053977	Sequence 1729	U18197
Sequence 1678	D88532	Sequence 1730	Y14738
Sequence 1679	AF007216	Sequence 1731	AF159056
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Sequence 1681	U24105	Sequence 1733	X04297
Sequence 1682	AB002387	Sequence 1734	AB011165
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Sequence 1684	AL117666	Sequence 1736	X15949
Sequence 1685	AL049951	Sequence 1737	D84105
Sequence 1686	L07540	Sequence 1738	AF125096
Sequence 1687	AL110242	Sequence 1739	AF053551
Sequence 1688	AF078861	Sequence 1740	AF007149
Sequence 1689	AF153612	Sequence 1741	AJ224326
Sequence 1690	M17254	Sequence 1742	M55543
Sequence 1691	D86967	Sequence 1743	AB032948
Sequence 1692	AB014548	Sequence 1744	AF091076
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Sequence 1694	AF022229	Sequence 1746	L13977
Sequence 1695	AL050051	Sequence 1747	M87284
Sequence 1698	AB033071	Sequence 1748	AB002382
Sequence 1697	AF052093	Sequence 1749	Z82022
Sequence 1698	Z13009	Sequence 1750	AF151908
Sequence 1699	AF156965	Sequence 1751	AL050107
Sequence 1700	AB018334	Sequence 1752	AF027824
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Sequence 1702	U17104	Sequence 1754	AL080201
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Sequence 1704	AL117567	Sequence 1756	D13119
Sequence 1705	AL080097	Sequence 1757	AF146277
Sequence 1706	AF151840	Sequence 1758	AF170583
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Sequence 1711	AF167160	Sequence 1763	AL110214
Sequence 1712	AB033091	Sequence 1764	AL080088
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Sequence 1714	M22918	Sequence 1766	AF078860
Sequence 1715	L05186	Sequence 1767	U03886

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Sequence 1768	L07033	Sequence 1820	D38555
Sequence 1769	U07919	Sequence 1821	D28540
Sequence 1770	D13629	Sequence 1822	AF191298
Sequence 1771	AF191018	Sequence 1823	AF042346
Sequence 1772	X04697	Sequence 1824	E01956
Sequence 1773	X81109	Sequence 1825	X70394
Sequence 1774	AL079298	Sequence 1826	U41060
Sequence 1775	X56958	Sequence 1827	AB000887
Sequence 1776	AL110126	Sequence 1828	U25182
Sequence 1777	X63679	Sequence 1829	X87241
Sequence 1778	X17620	Sequence 1830	AB007972
Sequence 1779	A21185	Sequence 1831	M63573
Sequence 1780	AF087942	Sequence 1832	X07270
Sequence 1781	E01954	Sequence 1833	AF124439
Sequence 1782	AF077200	Sequence 1834	D50406
Sequence 1783	D87078	Sequence 1835	J04443
Sequence 1784	AL050272	Sequence 1836	D21260
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Sequence 1786	U90144	Sequence 1838	D89092
Sequence 1787	D14696	Sequence 1839	K03002
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Sequence 1789	AF085845	Sequence 1841	M83822
Sequence 1790	AB033114	Sequence 1842	M55409
Sequence 1791	E01497	Sequence 1843	AB007885
Sequence 1792	AB007510	Sequence 1844	AL050198
Sequence 1793	D8 <b>7666</b>	Sequence 1845	M26481
Sequence 1794	D38524	Sequence 1846	M27544
Sequence 1795	E00882	Sequence 1847	X83544
Sequence 1798	X02875	Sequence 1848	D14812
Sequence 1797	AF0 <b>85224</b>	Sequence 1849	AF056433
Sequence 1798	AF071076	Sequence 1850	Z23064
Sequence 1799	AF102850	Sequence 1851	AB032983
Sequence 1800	AB <b>00236</b> 9	Sequence 1852	AL117514
Sequence 1801	AB011004	Sequence 1853	X59131
Sequence 1802	AL050289	Sequence 1854	AF073475
Sequence 1803	S41 <b>458</b>	Sequence 1855	AF044954
Sequence 1804	AF0880C4	Sequence 1856	X60221
Sequence 1805	AF086431	Sequence 1857	AB007882
Sequence 1806	AF010233	Sequence 1858	AF082513
Sequence 1807	AF086336	Sequence 1859	AF071202
Sequence 1808	AF057705	Sequence 1860	M26383
Sequence 1809	AB018280	Sequence 1861	X57351
Sequence 1810	AF016496	Sequence 1862	AF007142
Sequence 1811	M26663	Sequence 1863	AF020202
Sequence 1812	M17886	Sequence 1864	AB020686
Sequence 1813	X07466	Sequence 1865	X91257
Sequence 1814	D31767	Sequence 1866	D32051
Sequence 1815	L05779	Sequence 1867	AF147339
Sequence 1816	U07809	Sequence 1868	AB002306
Sequence 1817	AF137334	Sequence 1869	AF100756
Sequence 1818	AF056322	Sequence 1870	U01925
Sequence 1819	AB001636	Sequence 1871	D14878

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Sequence 1872	M22590	Sequence 1924	X27262
Sequence 1873	M93651	Sequence 1925	T59 <b>274</b>
Sequence 1874	AF054838	Sequence 1926	V69358
Sequence 1875	M69066	Sequence 1927	X989 <b>98</b>
Sequence 1876	AF017269	Sequence 1928	Z10658
Sequence 1877	J03007	Sequence 1929	X7657 <b>8</b>
Sequence 1878	AF019214	Sequence 1930	Z06652
Sequence 1879	AF039023	Sequence 1931	T37414
Sequence 1880	X66276	Sequence 1932	V56670
Sequence 1881	AL110179	Sequence 1933	V58682
Sequence 1882	D86984	Sequence 1934	Z15583
Sequence 1883	D90226	Sequence 1935	T18813
Sequence 1884	AL035304	Sequence 1936	X22116
Sequence 1885	D13643	Sequence 1937	T61701
Sequence 1886	K01911	Sequence 1938	Z00842
Sequence 1887	L28997	Sequence 1939	∨58 <b>756</b>
Sequence 1888	AB023173	Sequence 1940	V57903
Sequence 1889	Y00971	Sequence 1941	X03841
Sequence 1890	AB033899	Sequence 1942	Z17335
Sequence 1891	U90550	Sequence 1943	X26850
Sequence 1892	M10119	Sequence 1944	Z33658
Sequence 1893	AL133078	Sequence 1945	Z14392
Sequence 1894	J04607	Sequence 1946	V63172
Sequence 1895	J02642	Sequence 1947	Z15729
Sequence 1896	A06919	Sequence 1948	V34236
Sequence 1897	M34181	Sequ <b>ence 1949</b>	AC31081
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Sequence 1900	AF083441	Sequence 1952	AC33865
Sequence 1901	AF107405	Sequence 1953	AC36153
Sequence 1902	AF201077	Sequence 1954	AC29851
Sequence 1903	U02680	Sequence 1955	AC36206
Sequence 1904	L08599	Sequence 1956	AC36220
Sequence 1905	AF019226	Sequence 1957	AC36078
Sequence 1906	AF176574	Sequence 1958	AC36205
Sequence 1907	AB028977	Sequence 1959	AC35218
Sequence 1908	AJ000881	Sequence 1960	AC36180
Sequence 1909	D89678	Sequence 1961	AC36187
Sequence 1910	D50405	Sequence 1962	AC36079
Sequence 1911	D29641	Sequence 1963	AC36823
Sequence 1912	M34840	Sequence 1964	AC36709
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Sequence 1914	V62430	Sequence 1966	AA308091
Sequence 1915		Sequence 1967	AA442287
Sequence 1916		Sequence 1968	AA460440
Sequence 1917		Sequence 1969	AA513597
Sequence 1918		Sequence 1970	AA576425
Sequence 1919		Sequence 1971	AB032983
Sequence 1920		Sequence 1972	AF047020
Sequence 1921		Sequence 1973	AF103907
Sequence 1922		Sequence 1974	Al080439
Sequence 1923		Sequence 1975	A1088633
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Table 4-1

Sequence 1976	AI267321	Sequence 2028	AA663192
Sequence 1977	A1274700	Sequence 2029	AA449889
Sequence 1978	AI557336	Sequence 2030	AA209369
Sequence 1979	AI825988	Sequence 2031	AA354730
Sequence 1980	AL037105	Sequence 2032	AA354196
Sequence 1981	AW138181	Sequence 2033	Al791370
Sequence 1982	D14043	Sequence 2034	AA447433
Sequence 1983	L38961	Sequence 2035	N44743
Sequence 1984	M26880	Sequence 2038	AA528104
Sequence 1985	X77548	Sequence 2037	AA853882
Sequence 1986	AB001636	Sequence 2038	AA134726
Sequence 1987	AB008681	Sequence 2039	AA811333
Sequence 1988	AB020866	Sequence 2040	AL121340
Sequence 1989	AC000399	Sequence 2041	AA310201
Sequence 1990	AC002086	Sequence 2042	AI694572
Sequence 1991	AC003991	Sequence 2043	Al140921
Sequence 1992	AC004507	Sequence 2044	
Sequence 1993	AC004816	Sequence 2045	AA149579
Sequence 1994	AC005383	Sequence 2046	AA506729
Sequence 1995	AC005480	_ ·	AA877460
Sequence 1996	AC005538	Sequence 2047 Sequence 2048	AI199094
Sequence 1997	AC005831		AA975275
Sequence 1998	AC006153	Sequence 2049	N25122
Sequence 1999	AL031777	Sequence 2050	AI636911
Sequence 2000	AL117340	Sequence 2051	Al219803
Sequence 2001	AL121603	Sequence 2052	AA483044
Sequence 2002	AL121654	Sequence 2053	AA522552
Sequence 2003	AL122007	Sequence 2054	AI564822
Sequence 2004	AL135879	Sequence 2055	R63251
Sequence 2005	AP000303	Sequence 2056	AA526369
Sequence 2006	M11548	Sequence 2057	A1074652
Sequence 2007	S65371	Sequence 2058	AA603861
Sequence 2008	U13800	Sequence 2059	AA253339
Sequence 2009	U91321	Sequence 2060	AA313170
Sequence 2010	Z63584	Sequence 2061	AA228940
Sequence 2011	Z73900	Sequence 2062	AF001542
Sequence 2012	AA079188	Sequence 2063	A1090642
Sequence 2013	Al267838	Sequence 2064	AA035559
Sequence 2014	AA507791	Sequence 2065	AA304845
Sequence 2015	AA640474	Sequence 2066	AI652112
Sequence 2016	AI023483	Sequence 2067	AA058381
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Sequence 2018	AA325789	Sequence 2069	AA588023
Sequence 2019	AI151458	Sequence 2070	AI034131
Sequence 2020	AA393870	Sequence 2071	AA577463
	-	Sequence 2072	AW151704
Sequence 2021 Sequence 2022	H70269	Sequence 2073	A1267285
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Sequence 2024	AL043205	Sequence 2076	AL043046
Sequence 2025	AA056482	Sequence 2077	AI373120
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Sequence 2027	R69940	Sequence 2079	AA149337

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Sequence 2080	AA622482	Sequence 2132	AI269205
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Sequence 2082	Al453181	Sequence 2134	AA336412
Sequence 2083	AL121262	Sequence 2135	AW148750
Sequence 2084	AA306322	Sequence 2136	AA304846
Sequence 2085	Al635209	Sequence 2137	AA190998
Sequence 2086	AA973620	Sequence 2138	AA564505
Sequence 2087	AA323181	Sequence 2139	AI086059
Sequence 2088	AA399177	Sequence 2140	AA235224
Sequence 2089	R33609	Sequence 2141	AA405663
Sequence 2090	AW149706	Sequence 2142	C21370
Sequence 2091	AA878395	Sequence 2143	Al253335
Sequence 2092	Al198859	Sequence 2144	AA437224
Sequence 2093	AA099399	Sequence 2145	AA314893
Sequence 2094	AA467869	Sequence 2146	AI630616
Sequence 2095	AA7479 <b>0</b> 9	Sequence 2147	W27229
Sequence 2096	AA483008	Sequence 2148	AA195525
Sequence 2097	N51806	Sequence 2149	Al868421
Sequence 2098	AA548764	Sequence 2150	AA305969
Sequence 2099	AA229495	Sequence 2151	A1476310
Sequence 2100	AL046322	Sequence 2152	AA843845
Sequence 2101	AA146981	Sequence 2153	AI204916
Sequence 2102	AA152337	Sequence 2154	AA172043
Sequence 2103	AA280917	Sequence 2155	AI557235
Sequence 2104	AL043257 `	Sequence 2156	AA020899
Sequence 2105	AA026149	Sequence 2157	A1954167
Sequence 2106	AA732538	Sequence 2158	AI380932
Sequence 2107	AA876439	Sequence 2159	AI267502
Sequence 2108	AA405535	Sequence 2160	H30141
Sequence 2109	H80936	Sequence 2161	AA634408
Sequence 2110	AI378290	Sequence 2162	AA031287
Sequence 2111	AI761098	Sequence 2163	AA233109
Sequence 2112	AA188875	Sequence 2164	AA150421
Sequence 2113	AA703396	Sequence 2165	AA088714
Sequence 2114	AA312076	Sequence 2166	AA523354
Sequence 2115	AA424428	Sequence 2167	AI809963
Sequence 2116	T68102	Sequence 2168	AA130302
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Sequence 2122	AA159272	Sequence 2174	AA376424
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Sequence 2126	H25806	Sequence 2178	Al125653
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Sequence 2128	N52271	Sequence 2180	AA016281
Sequence 2129	AA046581	Sequence 2181	AL038234
Sequence 2130	AA579026	Sequence 2182	H08621
Sequence 2131	AA764981	Sequence 2183	AA703816

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Sequence 2184	AA487754	Sequence 2236	AA531227
Sequence 2185	Al217694	Sequence 2237	AA402408
Sequence 2186	Al671394	Sequ <b>ence 2238</b>	AA132904
Sequence 2187	AA516531	Sequence 2239	Al310138
Sequence 2188	AA322901	Sequence 2240	Al267216
Sequence 2189	AA188140	Sequence 2241	AA180941
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Sequence 2191	AA599324	Sequence 2243	AA504490
Sequence 2192	H08959	Sequence 2244	AA150220
Sequence 2193	AA308316	Sequence 2245	AA047021
Sequence 2194	N23606	Sequence 2246	AA450051
Sequence 2195	Al620796	Sequence 2247	H78478
Sequence 2196	AL043048	Sequence 2248	AA225115
Sequence 2197	H02308	Sequence 2249	C04910
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Sequence 2199	AA186586	Sequence 2251	N56690
Sequence 2200	Al697815	Sequence 2252	AA992882
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Sequence 2202	AA678072	Sequence 2254	AA306787
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Sequence 2204	AW177204	Sequence 2256	AA576425
Sequence 2205	AL045638	Sequence 2257	AA634120
Sequence 2206	N21237	Sequence 2258	AA675892
Sequence 2207	AA157008	Sequence 2259	AA056667
Sequence 2208	AA150672	Sequence 2260	A1087300
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Sequence 2221	AA361640	Sequence 2273	AA664480
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Sequence 2223	AA039967	Sequence 2275	AA843176
Sequence 2224	AA476586	Sequence 2276	Al216969
Sequence 2225	AA147100	Sequence 2277	AA313369
Sequence 2226	AI267612	Sequ <b>ence</b> 2278	AI217172
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Sequence 2233	AA534246	Sequence 2285	AA329562
Sequence 2234	AA205647	Sequence 2286	AA442517
Sequence 2235	AA351504	Sequence 2287	AL044723

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Sequence 2290	AA286678		AA449940
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Sequence 2292	AA426229	Ooquomee 20 / /	AA194162
Sequence 2293	AA527805	Ocquonio III io	AA244003
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Sequence 2295	AA314146	00400	AA701467
Sequence 2296	AA190408	Sequence 2348	AA128387
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Sequence 2321	AA371964	Sequence 2373	AA083370
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Sequence 255	7.1.41000	•	

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Sequence 2394	AA193086	Sequence 2446	A155 <b>7336</b>
Sequence 2395	AA180137	Sequence 2447	AA459 <b>455</b>
Sequence 2396	A1458673	Sequence 2448	AA476360
Sequence 2397	AA101742	Sequence 2449	AI080272
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Sequence 2399	AA865185	Sequence 2451	H03785
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-5400110 <b>6 2443</b>	747707071	Sequence 2495	VEC-0204

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Sequence 2600	A15 <b>57599</b>	Sequence 2652	AL039293
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Sequence 2602	AA431664	Sequence 2654	AA531563
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Sequence 2610	AA632742	Sequence 2662	AA352938
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Sequence 2640	AA594540	Sequence 2692	AA120781
Sequence 2641	AA701667	Sequence 2693	AA480844
Sequence 2642	AA147871	Sequence 2694	AA417097
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Sequence 2722	AA345189	Sequence 2774	AA563986
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Sequence 2733	AA036823	Sequence 2785	AA131961
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Sequence 2740	AA355264	Sequence 2792 Sequence 2793	Ai025519
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Sequence 2742	AA143034	Sequence 2795	AA353655
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Sequence 2749		Sequence 2802	AA070008
Sequence 2750		Sequence 2803	AA833759
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Sequence 2754		Sequence 2806	
Sequence 275	5 Al0855 <b>54</b>	Sequence 2807	FV (2.1017)

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Sequence 2812	AA446943	Sequence 2864	A1300489
Sequence 2813	AA356069	Sequence 2865	AA129 <b>574</b>
Sequence 2814	AI761190	Sequence 2866	AA577672
Sequence 2815	AA040694	Sequence 2867	AI762634
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Sequence 2819	AA427459	Sequence 2871	AA463533
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Sequence 2821	AA379449	Sequence 2873	AA206675
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Sequence 2823	AA333300	Sequence 2875	AA639972
Sequence 2824	AI253300	Sequence 2876	A1620284
Sequence 2825	Al253424	Sequence 2877	AA515908
Sequence 2826	Al374628	Sequence 2878	AA133742
Sequence 2827	AA069845	Sequence 2879	H49828
Sequence 2828	N86927	Sequence 2880	AA911245
Sequence 2829	AA469199	Sequence 2881	AA505816
Sequence 2830	AI267416	Sequence 2882	AA121836
Sequence 2831	AA234135	Sequence 2883	R52729
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Sequence 2833	A1808505	Sequence 2885	AA731537
Sequence 2834	H78888	Sequence 2886	AI588087
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Sequence 2837	AA812581	Sequence 2889	N24284
Sequence 2838	AA565214	Sequence 2890	AA088606
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Sequence 2840	AA534644	Sequence 2892	AA492280
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Sequence 2842	T84605	Sequence 2894	AA836834
Sequence 2843	AA523902	Sequence 2895	AA737421
Sequence 2844	AL037812	Sequence 2896	AA812081
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Sequence 2847	AA813266	Sequence 2899	AA362826
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Sequence 2851	AA836233	Sequence 2903	AA027126
Sequence 2852	AA557321	Sequence 2904	W63 <b>659</b>
Sequence 2853	Al198311	Sequence 2905	A1804327
Sequence 2854	AA903139	Sequence 2906	AA165 <b>152</b>
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Sequence 2856	AA323813	Sequence 2908	Al114489
Sequence 2857	A1469767	Sequence 2909	AW173278
Sequence 2858 Sequence 2859	R18975	Sequence 2910	AA296 <b>298</b>
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Table 4-1

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Sequence 2914 AA206004 Sequence 2966 AI688573 Sequence 2915 AI627750 Sequence 2967 AA227579 Sequence 2916 AI340246 Sequence 2966 AI340246 Sequence 2969 AA2375789 Sequence 2917 AA084197 Sequence 2918 AI267492 Sequence 2970 AA575898 AI267492 Sequence 2971 AA375898 Sequence 2919 AI207515 Sequence 2971 AA613916 Sequence 2921 AX56562 Sequence 2973 AA613916 Sequence 2921 W58562 Sequence 2973 AA613916 Sequence 2922 AA5151796 Sequence 2975 AA622756 Sequence 2924 AA312691 Sequence 2976 AA459243 Sequence 2926 AA312691 Sequence 2976 AA459433 Sequence 2926 AA554621 Sequence 2977 AA63943 Sequence 2978 AI30598 Sequence 2978 AA654621 Sequence 2978 AI30598 Sequence 2978 Sequence 2978 AI315498 Sequence 2978 AI315498 Sequence 2978 AI315498 Sequence 2979 AA513056 Sequence 2920 AI315498 Sequence 2921 AI26658 Sequence 2927 AI315498 Sequence 2928 AI272941 Sequence 2928 Sequence 2929 AI815498 Sequence 2929 AI3590878 Sequence 2929 AI3590878 Sequence 2929 AI3590878 Sequence 2931 AA262537 Sequence 2935 Sequence 2936 AA541515 Sequence 2936 AA541515 Sequence 2936 AA454921 Sequence 2936 AA541515 Sequence 2937 AA544983 Sequence 2938 AA321703 Sequence 2938 AA321703 Sequence 2939 AA564513 Sequence 2939 AA259189 Sequence 2940 AA564513 Sequence 2939 AA26503 Sequence 2941 AI8689759 Sequence 2939 AA263065 Sequence 2941 AI6666 Sequence 2944 AA59118 Sequence 2945 AA669818 Sequence 2946 AA3949118 Sequence 2949 AA669818 Sequence 2940 AA669818 Sequence 2940 AA669818 Sequence 2940 AA669818 Sequence 2940 AA66981	Sequence 2912	AA369786		
Sequence 2915 Al627750 Sequence 2967 AA227579 Sequence 2916 Al340246 Sequence 2968 AA131834 Sequence 2917 AD84197 Sequence 2968 AA131834 Sequence 2918 Al267492 Sequence 2970 AA575898 Sequence 2919 Al267492 Sequence 2971 AA195259 Sequence 2921 Al267492 Sequence 2971 AA195259 Sequence 2921 Miss6667 Sequence 2972 AA613916 Sequence 2921 W58562 Sequence 2973 Al815495 Sequence 2922 AA151796 Sequence 2974 AA452200 Sequence 2923 AV27968 Sequence 2975 AA459235 Sequence 2924 AA312691 Sequence 2977 AA452200 Sequence 2924 AA312691 Sequence 2977 AA452200 Sequence 2926 AA32698 Sequence 2977 AA452200 Sequence 2927 Al33698 Sequence 2977 AA452200 Sequence 2928 Al27691 Sequence 2977 Sequence 2978 AA594331 Sequence 2928 Al272941 Sequence 2978 T99603 Sequence 2929 AA15498 Sequence 2980 R39446 Sequence 2929 Al815498 Sequence 2980 R39446 Sequence 2930 H57064 Sequence 2981 AL037583 Sequence 2931 AA262537 Sequence 2984 Al267454 Sequence 2931 AA262537 Sequence 2984 Al74090 Sequence 2932 AL590878 Sequence 2984 Al74090 Sequence 2933 AA229872 Sequence 2986 AA7478931 Sequence 2934 Al815196 Sequence 2986 AA7478931 Sequence 2935 AA541515 Sequence 2986 AA7478931 Sequence 2936 AA454921 Sequence 2989 AA890010 Sequence 2937 AA164983 Sequence 2989 AA890010 Sequence 2938 AA034255 Sequence 2998 AA890010 Sequence 2940 AA564513 Sequence 2999 AA890010 Sequence 2941 AA364513 Sequence 2999 AA890010 Sequence 2944 AA349118 Sequence 2994 AA329118 Sequence 2994 AA32911 Sequence 2995 Sequence 2994 Sequence 2994 AA32911 Sequence 2995 Sequence 2995 Sequence 2995 Sequence 2995 Sequence 2995 Sequence 2996 Sequence 2996 Sequence 2996 Sequence 2996 Sequence 2997 Sequence 2998 Sequence 2998 Sequence 2999	Sequence 2913	AI394364	334233	
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Table 4-1

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